

Supplementary Table 1: Comparison of all the genes that are differentially regulated in the 18 transcriptome studies. The list of open reading frames (ORFs) are divided into core and accessory gene set as defined in Mathee *et al.* [PNAS. 2008.105(8). 3100-05].

| ORF             | Product name   | Function class   |  |  |  |
|-----------------|--|--|--|--|--|
| Transcriptic    | nal core set ORFs present in the P. aeruginosa core genome     |  |  |  |  |
| PA0044          | exoenzyme T  | Secreted Factors (toxins, enzymes, alginate)             |  |  |  |
| PA0045          | hypothetical protein   | Hypothetical, unclassified, unknown                      |  |  |  |
| PA0059          | osmotically inducible protein OsmC                             | Adaptation, Protection                                   |  |  |  |
| A0070           | hypothetical protein   | Membrane proteins  |  |  |  |
| A0078           | hypothetical protein   | Hypothetical, unclassified, unknown                      |  |  |  |
| A0082           | hypothetical protein   | Hypothetical, unclassified, unknown                      |  |  |  |
| PA0084          | conserved hypothetical protein                                 | Hypothetical, unclassified, unknown                      |  |  |  |
| A0084           | conserved hypothetical protein                                 | Hypothetical, unclassified, unknown                      |  |  |  |
| PA0140          | alkyl hydroperoxide reductase subunit F                        | Adaptation, Protection                                   |  |  |  |
| A0140<br>A0141  | conserved hypothetical protein                                 | Hypothetical, unclassified, unknown                      |  |  |  |
|                 | histidine porin OpdC   | Membrane proteins  |  |  |  |
| PA0162          |  |  |  |  |  |
| PA0183          | arylsulfatase  | Central intermediary metabolism                          |  |  |  |
| PA0184          | probable ATP-binding component of ABC transporter              | Transport of small molecules                             |  |  |  |
| PA0185          | probable permease of ABC transporter                           | Membrane proteins  |  |  |  |
| PA0186          | probable binding protein component of ABC transporter          | Transport of small molecules                             |  |  |  |
| PA0229          | dicarboxylic acid transporter PcaT                             | Membrane proteins  |  |  |  |
| PA0284          | hypothetical protein   | Hypothetical, unclassified, unknown                      |  |  |  |
| PA0290          | hypothetical protein   | Hypothetical, unclassified, unknown                      |  |  |  |
| PA0291          | Anaerobically-induced outer membrane porin OprE precursor      | Membrane proteins  |  |  |  |
| PA0424          | multidrug resistance operon repressor MexR                     | Transcriptional regulators                               |  |  |  |
| PA0447          | glutaryl-CoA dehydrogenase                                     | Fatty acid and phospholipid metabolism                   |  |  |  |
| PA0460          | hypothetical protein   | Hypothetical, unclassified, unknown                      |  |  |  |
| PA0471          | probable transmembrane sensor                                  | Two-component regulatory systems                         |  |  |  |
| PA0472          | probable sigma-70 factor, ECF subfamily                        | Transcriptional regulators                               |  |  |  |
| PA0514          | heme d1 biosynthesis protein NirL                              | Hypothetical, unclassified, unknown                      |  |  |  |
| PA0517          | probable c-type cytochrome precursor                           | Biosynthesis of cofactors, prosthetic groups and carrier |  |  |  |
| PA0518          | cytochrome c-551 precursor                                     | Biosynthesis of cofactors, prosthetic groups and carrier |  |  |  |
| PA0519          | nitrite reductase precursor                                    | Energy metabolism  |  |  |  |
| PA0524          | nitric-oxide reductase subunit B                               | Energy metabolism  |  |  |  |
| PA0572          | hypothetical protein   | Hypothetical, unclassified, unknown                      |  |  |  |
| PA0604          | probable binding protein component of ABC transporter          | Transport of small molecules                             |  |  |  |
| PA0605          | probable permease of ABC transporter                           | Membrane proteins  |  |  |  |
| PA0672          | heme oxygenase   | Biosynthesis of cofactors, prosthetic groups and carrier |  |  |  |
| PA0688          | probable binding protein component of ABC transporter          | Transport of small molecules                             |  |  |  |
| PA0762          | sigma factor AlgU  | Transcriptional regulators                               |  |  |  |
| PA0763          | anti-sigma factor MucA   | Transcriptional regulators                               |  |  |  |
| PA0764          | negative regulator for alginate biosynthesis MucB              | Transcriptional regulators                               |  |  |  |
| PA0775          | conserved hypothetical protein                                 | Hypothetical, unclassified, unknown                      |  |  |  |
| PA0848          | probable alkyl hydroperoxide reductase                         | Adaptation, Protection                                   |  |  |  |
| A0848<br>PA0849 | thioredoxin reductase 2  | Nucleotide biosynthesis and metabolism                   |  |  |  |
|                 |  | -  |  |  |  |
| PA0852          | chitin-binding protein CbpD precursor                          | Secreted Factors (toxins, enzymes, alginate)             |  |  |  |
| PA0866          | aromatic amino acid transport protein AroP2                    | Transport of small molecules                             |  |  |  |
| PA0874          | hypothetical protein   | Hypothetical, unclassified, unknown                      |  |  |  |
| PA0929          | two-component response regulator                               | Transport of small molecules                             |  |  |  |
| PA0938          | hypothetical protein   | Hypothetical, unclassified, unknown                      |  |  |  |
| PA0958          | Outer membrane porin OprD precursor                            | Transport of small molecules                             |  |  |  |
| PA0962          | probable dna-binding stress protein                            | Adaptation, Protection                                   |  |  |  |
| PA1071          | branched-chain amino acid transport protein BraF               | Transport of small molecules                             |  |  |  |
| PA1156          | NrdA, catalytic component of class Ia ribonucleotide reductase | Nucleotide biosynthesis and metabolism                   |  |  |  |
| PA1179          | two-component response regulator PhoP                          | Transcriptional regulators                               |  |  |  |
| PA1180          | two-component sensor PhoQ                                      | Two-component regulatory systems                         |  |  |  |
| PA1230          | hypothetical protein   | Membrane proteins  |  |  |  |
| PA1245          | hypothetical protein   | Membrane proteins  |  |  |  |
| PA1246          | alkaline protease secretion protein AprD                       | Secreted Factors (toxins, enzymes, alginate)             |  |  |  |
|                 | alkaline metalloproteinase precursor                           | Secreted Factors (toxins, enzymes, alginate)             |  |  |  |

| ORF                        | Product name   | Function class  |  |  |  |  |
|----------------------------|--|---|--|--|--|--|
| PA1289                     | hypothetical protein   | Hypothetical, unclassified, unknown                       |  |  |  |  |
| PA1300                     | probable sigma-70 factor, ECF subfamily  | Transcriptional regulators                                |  |  |  |  |
| PA1317                     | cytochrome o ubiquinol oxidase subunit II  | Energy metabolism   |  |  |  |  |
| PA1318                     | cytochrome o ubiquinol oxidase subunit I   | Energy metabolism   |  |  |  |  |
| PA1319                     | cytochrome o ubiquinol oxidase subunit II  | Energy metabolism   |  |  |  |  |
| PA1323                     | hypothetical protein   | Hypothetical, unclassified, unknown                       |  |  |  |  |
| PA1324                     | hypothetical protein   | Hypothetical, unclassified, unknown                       |  |  |  |  |
| PA1346                     | hypothetical protein   | Hypothetical, unclassified, unknown                       |  |  |  |  |
| PA1408                     | hypothetical protein   | Membrane proteins   |  |  |  |  |
| PA1541                     | probable drug efflux transporter   | Membrane proteins   |  |  |  |  |
| PA1544                     | transcriptional regulator Anr  | Transcriptional regulators                                |  |  |  |  |
| PA1546                     | oxygen-independent coproporphyrinogen III oxidase                                    | Biosynthesis of cofactors, prosthetic groups and carriers |  |  |  |  |
| PA1555                     | probable cytochrome c  | Energy metabolism   |  |  |  |  |
| PA1556                     | probable cytochrome c oxidase subunit  | Energy metabolism   |  |  |  |  |
| PA1557                     | probable cytochrome oxidase subunit (cbb3-type)                                      | Energy metabolism   |  |  |  |  |
| PA1657                     | conserved hypothetical protein   | Hypothetical, unclassified, unknown                       |  |  |  |  |
| PA1673                     | hypothetical protein   | Hypothetical, unclassified, unknown                       |  |  |  |  |
| PA1691                     | translocation protein in type III secretion  | Protein secretion/export apparatus                        |  |  |  |  |
| PA1696                     | translocation protein in type III secretion  | Protein secretion/export apparatus                        |  |  |  |  |
| PA1698                     | Type III secretion outer membrane protein PopN precursor                             | Membrane proteins   |  |  |  |  |
| PA1699                     | conserved hypothetical protein in type III secretion                                 | Protein secretion/export apparatus                        |  |  |  |  |
| PA1700                     | conserved hypothetical protein in type III secretion                                 | Protein secretion/export apparatus                        |  |  |  |  |
| PA1701                     | conserved hypothetical protein in type III secretion                                 | Protein secretion/export apparatus                        |  |  |  |  |
| PA1705                     | regulator in type III secretion  | Protein secretion/export apparatus                        |  |  |  |  |
| PA1706                     | type III secretion protein PcrV  | Protein secretion/export apparatus                        |  |  |  |  |
| PA1707                     | regulatory protein PcrH  | Secreted Factors (toxins, enzymes, alginate)              |  |  |  |  |
| PA1708                     | translocator protein PopB  | Protein secretion/export apparatus                        |  |  |  |  |
| PA1709                     | Translocator outer membrane protein PopD precursor                                   | Protein secretion/export apparatus                        |  |  |  |  |
| PA1710                     | ExsC, exoenzyme S synthesis protein C precursor.                                     | Translation, post-translational modification, degradation |  |  |  |  |
| PA1711                     | ExsE   | Hypothetical, unclassified, unknown                       |  |  |  |  |
| PA1712                     | exoenzyme S synthesis protein B  | Translation, post-translational modification, degradation |  |  |  |  |
| PA1713                     | transcriptional regulator ExsA   | Protein secretion/export apparatus                        |  |  |  |  |
| PA1714                     | ExsD   | Hypothetical, unclassified, unknown                       |  |  |  |  |
| PA1715                     | type III export apparatus protein  | Protein secretion/export apparatus                        |  |  |  |  |
| PA1716                     | Type III secretion outer membrane protein PscC precursor                             | Protein secretion/export apparatus                        |  |  |  |  |
| PA1717                     | type III export protein PscD   | Protein secretion/export apparatus                        |  |  |  |  |
| PA1719                     | type III export protein PscF   | Protein secretion/export apparatus                        |  |  |  |  |
| PA1720                     | type III export protein PscG   | Protein secretion/export apparatus                        |  |  |  |  |
| PA1721                     | type III export protein PscH   | Protein secretion/export apparatus                        |  |  |  |  |
| PA1722                     | type III export protein PscI   | Protein secretion/export apparatus                        |  |  |  |  |
| PA1723                     | type III export protein PscJ   | Protein secretion/export apparatus                        |  |  |  |  |
| PA1761                     | hypothetical protein   | Hypothetical, unclassified, unknown                       |  |  |  |  |
| PA1793                     | peptidyl-prolyl cis-trans isomerase B  | Translation, post-translational modification, degradation |  |  |  |  |
| PA1869                     | probable acyl carrier protein  | Fatty acid and phospholipid metabolism                    |  |  |  |  |
| PA1894                     | hypothetical protein   | Hypothetical, unclassified, unknown                       |  |  |  |  |
| PA1902                     | phenazine biosynthesis protein PhzD  | Secreted Factors (toxins, enzymes, alginate)              |  |  |  |  |
| PA1904                     | probable phenazine biosynthesis protein  | Secreted Factors (toxins, enzymes, alginate)              |  |  |  |  |
| PA1982                     | quinoprotein alcohol dehydrogenase   | Carbon compound catabolism                                |  |  |  |  |
| PA2080                     | hypothetical protein   | Hypothetical, unclassified, unknown                       |  |  |  |  |
| PA2081                     | kynurenine formamidase, KynB   | Hypothetical, unclassified, unknown                       |  |  |  |  |
| PA2087                     | hypothetical protein   | Hypothetical, unclassified, unknown                       |  |  |  |  |
| PA2088                     | hypothetical protein   | Hypothetical, unclassified, unknown                       |  |  |  |  |
| PA2111                     | hypothetical protein   | Hypothetical, unclassified, unknown                       |  |  |  |  |
|                            | conserved hypothetical protein   | Hypothetical, unclassified, unknown                       |  |  |  |  |
| PAZITZ                     |  |   |  |  |  |  |
| PA2112<br>PA2113           | pyroglutamate porin OpdO   | Transport of small molecules                              |  |  |  |  |
| PA2112<br>PA2113<br>PA2114 | pyroglutamate porin OpdO<br>probable major facilitator superfamily (MFS) transporter | Transport of small molecules<br>Membrane proteins         |  |  |  |  |

| ORF              | Product name   | Function class                               |  |  |
|------------------|--|--|--|--|
| PA2146           | conserved hypothetical protein                                 | Hypothetical, unclassified, unknown          |  |  |
| PA2158           | probable alcohol dehydrogenase (Zn-dependent)                  | Putative enzymes                             |  |  |
| PA2159           | conserved hypothetical protein                                 | Hypothetical, unclassified, unknown          |  |  |
| PA2166           | hypothetical protein   | Hypothetical, unclassified, unknown          |  |  |
| PA2171           | hypothetical protein   | Hypothetical, unclassified, unknown          |  |  |
| PA2173           | hypothetical protein   | Hypothetical, unclassified, unknown          |  |  |
| PA2175           | probable sensor/response regulator hybrid                      | Two-component regulatory systems             |  |  |
| PA2177<br>PA2191 |  | Secreted Factors (toxins, enzymes, alginate) |  |  |
|                  | adenylate cyclase ExoY   |  |  |  |
| PA2193           | hydrogen cyanide synthase HcnA                                 | Central intermediary metabolism              |  |  |
| PA2194           | hydrogen cyanide synthase HcnB                                 | Central intermediary metabolism              |  |  |
| PA2195           | hydrogen cyanide synthase HcnC                                 | Central intermediary metabolism              |  |  |
| PA2247           | 2-oxoisovalerate dehydrogenase (alpha subunit)                 | Amino acid biosynthesis and metabolism       |  |  |
| PA2248           | 2-oxoisovalerate dehydrogenase (beta subunit)                  | Amino acid biosynthesis and metabolism       |  |  |
| PA2260           | hypothetical protein   | Hypothetical, unclassified, unknown          |  |  |
| PA2261           | probable 2-ketogluconate kinase                                | Carbon compound catabolism                   |  |  |
| PA2262           | probable 2-ketogluconate transporter                           | Membrane proteins                            |  |  |
| PA2263           | probable 2-hydroxyacid dehydrogenase                           | Putative enzymes                             |  |  |
| PA2264           | conserved hypothetical protein                                 | Hypothetical, unclassified, unknown          |  |  |
| PA2265           | gluconate dehydrogenase  | Carbon compound catabolism                   |  |  |
| PA2266           | probable cytochrome c precursor                                | Carbon compound catabolism                   |  |  |
| PA2299           | probable transcriptional regulator                             | Transcriptional regulators                   |  |  |
| PA2302           | probable non-ribosomal peptide synthetase                      | Putative enzymes                             |  |  |
| PA2304           | hypothetical protein   | Hypothetical, unclassified, unknown          |  |  |
| PA2305           | probable non-ribosomal peptide synthetase                      | Putative enzymes                             |  |  |
|                  |  |  |  |  |
| PA2307           | probable permease of ABC transporter                           | Membrane proteins                            |  |  |
| PA2309           | hypothetical protein   | Hypothetical, unclassified, unknown          |  |  |
| PA2310           | hypothetical protein   | Putative enzymes                             |  |  |
| PA2331           | hypothetical protein   | Membrane proteins                            |  |  |
| PA2385           | PvdQ   | Adaptation, Protection                       |  |  |
| PA2386           | L-ornithine N5-oxygenase                                       | Adaptation, Protection                       |  |  |
| PA2389           | conserved hypothetical protein                                 | Hypothetical, unclassified, unknown          |  |  |
| PA2392           | PvdP   | Adaptation, Protection                       |  |  |
| PA2393           | probable dipeptidase precursor                                 | Central intermediary metabolism              |  |  |
| PA2394           | PvdN   | Adaptation, Protection                       |  |  |
| PA2396           | pyoverdine synthetase F  | Secreted Factors (toxins, enzymes, alginate) |  |  |
| PA2397           | pyoverdine biosynthesis protein PvdE                           | Membrane proteins                            |  |  |
| PA2403           | hypothetical protein   | Membrane proteins                            |  |  |
| PA2404           | hypothetical protein   | Membrane proteins                            |  |  |
| PA2405           | hypothetical protein   | Hypothetical, unclassified, unknown          |  |  |
| PA2407           | probable adhesion protein                                      | Motility & Attachment                        |  |  |
| PA2408           | probable ATP-binding component of ABC transporter              | Transport of small molecules                 |  |  |
| PA2400           | probable permease of ABC transporter                           | Membrane proteins                            |  |  |
|                  |  |  |  |  |
| PA2410           | hypothetical protein   | Hypothetical, unclassified, unknown          |  |  |
| PA2411           | probable thioesterase  | Adaptation, Protection                       |  |  |
| PA2412           | conserved hypothetical protein                                 | Hypothetical, unclassified, unknown          |  |  |
| PA2413           | L-2,4-diaminobutyrate:2-ketoglutarate 4-aminotransferase, PvdH | Adaptation, Protection                       |  |  |
| PA2414           | L-sorbosone dehydrogenase                                      | Carbon compound catabolism                   |  |  |
| PA2415           | hypothetical protein   | Membrane proteins                            |  |  |
| PA2424           | PvdL   | Adaptation, Protection                       |  |  |
| PA2425           | PvdG   | Adaptation, Protection                       |  |  |
| PA2426           | sigma factor PvdS  | Transcriptional regulators                   |  |  |
| PA2433           | hypothetical protein   | Hypothetical, unclassified, unknown          |  |  |
| PA2468           | ECF sigma factor FoxI  | Transcriptional regulators                   |  |  |
| PA2505           | tyrosine porin OpdT  | Transport of small molecules                 |  |  |
| PA2523           | probable two-component response regulator                      | Transcriptional regulators                   |  |  |
| PA2548           | hypothetical protein   | Hypothetical, unclassified, unknown          |  |  |
| PA2588           | probable transcriptional regulator                             | Transcriptional regulators                   |  |  |
| гадзөө           | probable transcriptional regulator                             | rranscriptional regulators                   |  |  |

| ORF Product name           |   | Function class  |  |  |  |
|----------------------------|---|---|--|--|--|
| PA2592                     | probable periplasmic spermidine/putrescine-binding protein    | Transport of small molecules                              |  |  |  |
| PA2630                     | conserved hypothetical protein                                | Hypothetical, unclassified, unknown                       |  |  |  |
| PA2634                     | isocitrate lyase  | Putative enzymes  |  |  |  |
| PA2687                     | two-component sensor PfeS                                     | Two-component regulatory systems                          |  |  |  |
| PA2747                     | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |  |  |
| PA2760                     | probable outer membrane protein precursor                     | Transport of small molecules                              |  |  |  |
| PA2808                     | two-component response repressor, PtrA                        | Transcriptional regulators                                |  |  |  |
|                            |   |   |  |  |  |
| PA2850                     | organic hydroperoxide resistance protein                      | Adaptation, Protection                                    |  |  |  |
| PA2896                     | probable sigma-70 factor, ECF subfamily                       | Transcriptional regulators                                |  |  |  |
| PA3235                     | conserved hypothetical protein                                | Membrane proteins   |  |  |  |
| PA3278                     | hypothetical protein  | Membrane proteins   |  |  |  |
| PA3281                     | hypothetical protein  | Membrane proteins   |  |  |  |
| PA3283                     | conserved hypothetical protein                                | Hypothetical, unclassified, unknown                       |  |  |  |
| PA3284                     | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |  |  |
| PA3287                     | conserved hypothetical protein                                | Hypothetical, unclassified, unknown                       |  |  |  |
| PA3328                     | probable FAD-dependent monooxygenase                          | Putative enzymes  |  |  |  |
| PA3369                     | hypothetical protein  | Membrane proteins   |  |  |  |
| PA3371                     | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |  |  |
| PA3413                     | conserved hypothetical protein                                | Hypothetical, unclassified, unknown                       |  |  |  |
| PA3414                     | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |  |  |
|                            |   |   |  |  |  |
| PA3416                     | probable pyruvate dehydrogenase E1 component, beta chain      | Energy metabolism   |  |  |  |
| PA3446                     | conserved hypothetical protein                                | Hypothetical, unclassified, unknown                       |  |  |  |
| PA3447                     | probable ATP-binding component of ABC transporter             | Transport of small molecules                              |  |  |  |
| PA3448                     | probable permease of ABC transporter                          | Membrane proteins   |  |  |  |
| PA3450                     | probable antioxidant protein                                  | Adaptation, Protection                                    |  |  |  |
| PA3460                     | probable acetyltransferase                                    | Putative enzymes  |  |  |  |
| PA3461                     | conserved hypothetical protein                                | Hypothetical, unclassified, unknown                       |  |  |  |
| PA3515                     | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |  |  |
| PA3516                     | probable lyase  | Putative enzymes  |  |  |  |
| PA3517                     | probable lyase  | Carbon compound catabolism                                |  |  |  |
| PA3518                     | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |  |  |
| PA3519                     | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |  |  |
| PA3520                     | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |  |  |
| PA3523                     | probable RND efflux membrane fusion protein precursor         | Transport of small molecules                              |  |  |  |
| PA3531                     | bacterioferritin  | Transport of small molecules                              |  |  |  |
|                            |   |   |  |  |  |
| PA3573                     | probable major facilitator superfamily (MFS) transporter      | Adaptation, Protection                                    |  |  |  |
| PA3600                     | conserved hypothetical protein                                | Translation, post-translational modification, degradation |  |  |  |
| PA3690                     | probable metal-transporting P-type ATPase                     | Membrane proteins   |  |  |  |
| PA3691                     | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |  |  |
| PA3709                     | probable major facilitator superfamily (MFS) transporter      | Membrane proteins   |  |  |  |
| PA3790                     | Putative copper transport outer membrane porin OprC precursor | Transport of small molecules                              |  |  |  |
| PA3842                     | probable chaperone  | Secreted Factors (toxins, enzymes, alginate)              |  |  |  |
| PA3899                     | probable sigma-70 factor, ECF subfamily                       | Transcriptional regulators                                |  |  |  |
| PA3900                     | probable transmembrane sensor                                 | Membrane proteins   |  |  |  |
| PA3904                     | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |  |  |
| PA3936                     | probable permease of ABC taurine transporter                  | Membrane proteins   |  |  |  |
| PA3937                     | probable ATP-binding component of ABC taurine transporter     | Transport of small molecules                              |  |  |  |
| PA3938                     | probable periplasmic taurine-binding protein precursor        | Transport of small molecules                              |  |  |  |
|                            |   |   |  |  |  |
| PA3954                     | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |  |  |
| PA4067                     | Outer membrane protein OprG precursor                         | Membrane proteins   |  |  |  |
| PA4139                     | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |  |  |
| PA4156                     | probable TonB-dependent receptor                              | Transport of small molecules                              |  |  |  |
| PA4168                     | second ferric pyoverdine receptor FpvB                        | Transport of small molecules                              |  |  |  |
| D4 4171                    | probable protease   | Putative enzymes  |  |  |  |
| PA41/1                     |   |   |  |  |  |
|                            | probable porin  | Membrane proteins   |  |  |  |
| PA4171<br>PA4179<br>PA4217 |   | Membrane proteins<br>Putative enzymes                     |  |  |  |

| ORF              | Product name   | Function class   |
|------------------|--|--|
| DA 4210          | have all address and sin   | Manufacture and dime   |
| PA4219           | hypothetical protein   | Membrane proteins  |
| PA4220<br>PA4221 | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA4221<br>PA4222 | Fe(III)-pyochelin outer membrane receptor precursor  | Transport of small molecules   |
| PA4222<br>PA4224 | probable ATP-binding component of ABC transporter  | Transport of small molecules   |
|                  | pyochelin biosynthetic protein PchG  | Transport of small molecules   |
| PA4225           | pyochelin synthetase<br>dihydroaeruginoic acid synthetase                                    | Secreted Factors (toxins, enzymes, alginate)   |
| PA4226           | transcriptional regulator PchR   | Secreted Factors (toxins, enzymes, alginate)   |
| PA4227<br>PA4228 | pyochelin biosynthesis protein PchD  | Transcriptional regulators   |
| PA4228<br>PA4229 | pyochelin biosynthesis protein PchD  | Secreted Factors (toxins, enzymes, alginate)   |
|                  |  | Secreted Factors (toxins, enzymes, alginate)   |
| PA4230           | salicylate biosynthesis protein PchB   | Secreted Factors (toxins, enzymes, alginate)   |
| PA4231           | salicylate biosynthesis isochorismate synthase   | Secreted Factors (toxins, enzymes, alginate)   |
| PA4296           | two-component response regulator, PprB   | Transcriptional regulators   |
| PA4359           | conserved hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA4429           | probable cytochrome c1 precursor   | Energy metabolism  |
| PA4430           | probable cytochrome b  | Energy metabolism  |
| PA4431           | probable iron-sulfur protein   | Putative enzymes   |
| PA4467           | hypothetical protein   | Membrane proteins  |
| PA4468           | superoxide dismutase   | Adaptation, Protection   |
| PA4469           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA4470           | fumarate hydratase   | Energy metabolism  |
| PA4471           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA4480           | rod shape-determining protein MreC   | Cell wall / LPS / capsule  |
| PA4501           | Glycine-glutamate dipeptide porin OpdP   | Transport of small molecules   |
| PA4563           | 30S ribosomal protein S20  | Central intermediary metabolism  |
| PA4570           | hypothetical protein<br>protein activator  | Hypothetical, unclassified, unknown  |
| PA4590           | catalase   | Transport of small molecules   |
| PA4613           |  | Adaptation, Protection   |
| PA4670<br>PA4673 | ribose-phosphate pyrophosphokinase   | Carbon compound catabolism   |
| PA4073<br>PA4708 | conserved hypothetical protein<br>Heme-transport protein, PhuT                               | Hypothetical, unclassified, unknown<br>Transport of small molecules                              |
|                  |  |  |
| PA4709           | probable hemin degrading factor  | Putative enzymes   |
| PA4710           | Haem/Haemoglobin uptake outer membrane receptor PhuR precursor                               | Transport of small molecules   |
| PA4719<br>PA4738 | probable transporter<br>conserved hypothetical protein                                       | Membrane proteins<br>Hypothetical, unclassified, unknown   |
|                  |  | Hypothetical, unclassified, unknown  |
| PA4739           | conserved hypothetical protein   | Membrane proteins  |
| PA4876<br>PA4880 | osmotically inducible lipoprotein OsmE   | -  |
| PA4880<br>PA4896 | probable bacterioferritin  | Central intermediary metabolism<br>Transcriptional regulators                                    |
| PA4896<br>PA4909 | probable sigma-70 factor, ECF subfamily<br>probable ATP-binding component of ABC transporter | Transport of small molecules   |
|                  |  |  |
| PA4981           | probable amino acid permease   | Membrane proteins  |
| PA5261           | alginate biosynthesis regulatory protein AlgR  | Secreted Factors (toxins, enzymes, alginate)   |
| PA5311           | probable major facilitator superfamily (MFS) transporter                                     | Membrane proteins  |
| PA5315           | 50S ribosomal protein L33  | Translation, post-translational modification, degradation<br>Hypothetical, unclassified, unknown |
| PA5407           | hypothetical protein   |  |
| PA5469           | conserved hypothetical protein   | Membrane proteins  |
| PA5482           | hypothetical protein   | Membrane proteins  |
| PA5530           | probable MFS dicarboxylate transporter   | Membrane proteins  |

# Transcriptional core set ORFs present in the *P. aeruginosa* PAO1 accessory genome

| PA0203 | probable binding protein component of ABC transporter | Transport of small molecules             |
|--------|---|--|
| PA0204 | probable permease of ABC transporter                  | Membrane proteins                        |
| PA0205 | probable permease of ABC transporter                  | Membrane proteins                        |
| PA0206 | probable ATP-binding component of ABC transporter     | Transport of small molecules             |
| PA0633 | hypothetical protein                                  | Related to phage, transposon, or plasmid |
| PA0691 | hypothetical protein                                  | Hypothetical, unclassified, unknown      |

| ORF    | Product name                                      | Function class                               |  |  |
|--------|---|--|--|--|
|        |   |  |  |  |
| PA1414 | hypothetical protein                              | Hypothetical, unclassified, unknown          |  |  |
| PA1592 | hypothetical protein                              | Hypothetical, unclassified, unknown          |  |  |
| PA1718 | type III export protein PscE                      | Protein secretion/export apparatus           |  |  |
| PA2033 | hypothetical protein                              | Hypothetical, unclassified, unknown          |  |  |
| PA2034 | hypothetical protein                              | Hypothetical, unclassified, unknown          |  |  |
| PA2184 | conserved hypothetical protein                    | Hypothetical, unclassified, unknown          |  |  |
| PA2190 | conserved hypothetical protein                    | Hypothetical, unclassified, unknown          |  |  |
| PA2308 | probable ATP-binding component of ABC transporter | Transport of small molecules                 |  |  |
| PA2398 | ferripyoverdine receptor                          | Transport of small molecules                 |  |  |
| PA2399 | pyoverdine synthetase D                           | Secreted Factors (toxins, enzymes, alginate) |  |  |
| PA2400 | PvdJ  | Adaptation, Protection                       |  |  |
| PA2402 | probable non-ribosomal peptide synthetase         | Putative enzymes                             |  |  |
| PA2452 | hypothetical protein                              | Hypothetical, unclassified, unknown          |  |  |
| PA3370 | hypothetical protein                              | Membrane proteins                            |  |  |
| PA3477 | transcriptional regulator RhlR                    | Adaptation, Protection                       |  |  |
| PA3841 | exoenzyme S                                       | Secreted Factors (toxins, enzymes, alginate) |  |  |
| PA3866 | pyocin protein                                    | Adaptation, Protection                       |  |  |
| PA3923 | hypothetical protein                              | Hypothetical, unclassified, unknown          |  |  |
| PA4193 | probable permease of ABC transporter              | Membrane proteins                            |  |  |
| PA4223 | probable ATP-binding component of ABC transporter | Membrane proteins                            |  |  |
| PA4306 | Type IVb pilin, Flp                               | Motility & Attachment                        |  |  |
| PA4555 | type 4 fimbrial biogenesis protein PilY2          | Motility & Attachment                        |  |  |
| PA4748 | triosephosphate isomerase                         | Central intermediary metabolism              |  |  |
|        |   |  |  |  |

Supplementary Table 2: Functional categorization of genes that are uniquely expressed under the different study conditions.

|      |   | Percentage of differentially regulated unique genes (minus the core set) |    |             |     |        |         |    |    |      |         |
|------|---|--|----|-------------|-----|--------|---------|----|----|------|---------|
| Code | Class   | Iron   | QS | Oxi. Stress | РСР | Copper | Sulfate |    |    | RsmA | Total % |
| a    | DNA replication, recombination, modification and repair   | 0  | 1  | 1           | 0   | 0      | 0       | 0  | 0  | 0    | 0.4     |
| b    | Fatty acid and phospholipid metabolism                    | 0  | 2  | 0           | 1   | 0      | 0       | 0  | 2  | 1    | 1.2     |
| c    | Hypothetical, unclassified, unknown                       | 27   | 32 | 26          | 31  | 14     | 14      | 3  | 40 | 1    | 29.7    |
| d    | Membrane proteins   | 22   | 12 | 8           | 11  | 33     | 14      | 0  | 12 | 8    | 13.3    |
| e    | Amino acid biosynthesis and metabolism                    | 3  | 3  | 2           | 4   | 0      | 0       | 3  | 2  | 6    | 2.5     |
| f    | Translation, post-translational modification, degradation | 1  | 1  | 3           | 5   | 2      | 0       | 3  | 4  | 1    | 2.4     |
| g    | Cell wall / LPS / capsule                                 | 1  | 1  | 1           | 3   | 0      | 0       | 0  | 5  | 6    | 2.2     |
| h    | Transport of small molecules                              | 10   | 7  | 9           | 12  | 19     | 17      | 0  | 4  | 8    | 7.7     |
| i    | Energy metabolism   | 4  | 3  | 15          | 3   | 7      | 3       | 0  | 3  | 7    | 5.1     |
| j    | Biosynthesis of cofactors, prosthetic groups and carriers | 4  | 2  | 2           | 3   | 1      | 0       | 0  | 0  | 3    | 1.7     |
| k    | Adaptation, Protection                                    | 3  | 2  | 4           | 0   | 5      | 0       | 3  | 3  | 14   | 3.1     |
| 1    | Transcriptional regulators                                | 7  | 6  | 3           | 7   | 10     | 3       | 37 | 5  | 3    | 5.8     |
| m    | Two-component regulatory systems                          | 1  | 1  | 0           | 0   | 4      | 0       | 10 | 2  | 1    | 1.1     |
| n    | Secreted Factors (toxins, enzymes, alginate)              | 7  | 2  | 0           | 1   | 1      | 0       | 0  | 1  | 7    | 2.2     |
| 0    | Putative enzymes  | 7  | 10 | 7           | 9   | 3      | 34      | 3  | 7  | 0    | 8.0     |
| р    | Chaperones & heat shock proteins                          | 1  | 0  | 0           | 1   | 0      | 0       | 0  | 0  | 0    | 0.2     |
| q    | Central intermediary metabolism                           | 1  | 3  | 3           | 4   | 0      | 0       | 0  | 1  | 0    | 2.1     |
| r    | Nucleotide biosynthesis and metabolism                    | 0  | 1  | 0           | 0   | 0      | 14      | 0  | 1  | 1    | 0.7     |
| S    | Carbon compound catabolism                                | 1  | 6  | 4           | 0   | 0      | 14      | 0  | 1  | 11   | 3.9     |
| t    | Motility & Attachment                                     | 1  | 0  | 1           | 0   | 0      | 0       | 0  | 2  | 13   | 1.2     |
| u    | Chemotaxis  | 1  | 0  | 1           | 1   | 0      | 0       | 0  | 1  | 0    | 0.4     |
| v    | Related to phage, transposon, or plasmid                  | 0  | 5  | 10          | 1   | 0      | 0       | 0  | 0  | 1    | 3.2     |
| w    | Non-coding RNA gene                                       | 0  | 0  | 0           | 0   | 0      | 0       | 0  | 0  | 0    | 0.1     |
| x    | Protein secretion/export apparatus                        | 1  | 1  | 0           | 0   | 0      | 0       | 33 | 1  | 1    | 1.1     |
| у    | Cell division   | 0  | 0  | 0           | 0   | 0      | 0       | 0  | 1  | 0    | 0.3     |
| zA   | Antibiotic resistance and susceptibility                  | 0  | 0  | 0           | 1   | 0      | 0       | 0  | 0  | 1    | 0.2     |
| zB   | Transcription, RNA processing and degradation             | 0  | 0  | 0           | 0   | 0      | 0       | 3  | 1  | 0    | 0.4     |

Note: QS- quorum sensing, Oxi. Stress- oxidative stress, PCP- pentachlorophenol stress

Supplementary Table 3: List of genes that are uniquely expressed in the various transcriptome studies. The various conditions include iron response, quorum sensing, oxidative stress, pentachlorophenol stress, copper stress, sulfate starvation, osmotic shock, biofilm formation, alginate production and loss of the post-transcriptional regulator RsmA.

Iron

| ORF              | Product name  | Function class   |
|------------------|---|--|
| PA0013           | conserved hypothetical protein  | Membrane proteins  |
| PA0027           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA0128           | conserved hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA0203           | probable binding protein component of ABC transporter   | Transport of small molecules   |
| PA0223           | probable dihydrodipicolinate synthetase   | Cell wall / LPS / capsule  |
| PA0224           | probable aldolase   | Putative enzymes   |
| PA0251           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA0252           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA0265           | succinate-semialdehyde dehydrogenase  | Central intermediary metabolism  |
| PA0266           | 4-aminobutyrate aminotransferase  | Central intermediary metabolism  |
| PA0296           | probable glutamine synthetase   | Putative enzymes   |
| PA0346           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA0417           | probable chemotaxis protein   | Chemotaxis   |
| PA0433<br>PA0460 | hypothetical protein<br>hypothetical protein  | Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown |
| PA0400<br>PA0470 | Ferrichrome receptor FiuA   | Transport of small molecules   |
| PA0470<br>PA0471 | probable transmembrane sensor   | Two-component regulatory systems   |
| PA0471           | probable sigma-70 factor, ECF subfamily   | Transcriptional regulators   |
| PA0500           | biotin synthase   | Biosynthesis of cofactors, prosthetic groups and carriers                  |
| PA0527           | transcriptional regulator Dnr   | Transcriptional regulators   |
| PA0539           | hypothetical protein  | Membrane proteins  |
| PA0605           | probable permease of ABC transporter  | Membrane proteins  |
| PA0672           | heme oxygenase  | Biosynthesis of cofactors, prosthetic groups and carriers                  |
| PA0707           | transcriptional regulator ToxR  | Transcriptional regulators   |
| PA0725           | hypothetical protein of bacteriophage Pf1   | Hypothetical, unclassified, unknown  |
| PA0774           | conserved hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA0801           | hypothetical protein  | Membrane proteins  |
| PA0802           | hypothetical protein  | Membrane proteins  |
| PA0818           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA0842           | probable glycosyl transferase   | Putative enzymes   |
| PA0844           | hemolytic phospholipase C precursor   | Secreted Factors (toxins, enzymes, alginate)                               |
| PA0848           | probable alkyl hydroperoxide reductase  | Adaptation, Protection   |
| PA0930           | two-component sensor  | Transport of small molecules   |
| PA0931           | ferric enterobactin receptor PirA   | Transport of small molecules   |
| PA0968           | conserved hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA1003           | Transcriptional regulator MvfR  | Biosynthesis of cofactors, prosthetic groups and carriers                  |
| PA1108           | probable major facilitator superfamily (MFS) transporter  | Membrane proteins  |
| PA1134           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA1148           | exotoxin A precursor  | Secreted Factors (toxins, enzymes, alginate)                               |
| PA1211           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA1247           | alkaline protease secretion protein AprE  | Secreted Factors (toxins, enzymes, alginate)                               |
| PA1248           | Alkaline protease secretion outer membrane protein AprF precursor                                       | Secreted Factors (toxins, enzymes, alginate)                               |
| PA1249           | alkaline metalloproteinase precursor  | Secreted Factors (toxins, enzymes, alginate)                               |
| PA1300           | probable sigma-70 factor, ECF subfamily   | Transcriptional regulators   |
| PA1301<br>PA1302 | probable transmembrane sensor   | Membrane proteins  |
| PA1302<br>PA1316 | probable heme utilization protein precursor<br>probable major facilitator superfamily (MFS) transporter | Membrane proteins<br>Membrane proteins                                     |
| PA1316<br>PA1317 | cytochrome o ubiquinol oxidase subunit II   | Energy metabolism  |
| PA1317<br>PA1318 | cytochrome o ubiquinol oxidase subunit I  | Energy metabolism  |
| PA1318           | cytochrome o ubiquinol oxidase subunit III  | Energy metabolism  |
| PA1320           | cytochrome o ubiquinol oxidase subunit IV   | Energy metabolism  |
| PA1320           | cytochrome o ubiquinol oxidase protein CyoE   | Energy metabolism  |
| PA1363           | probable sigma-70 factor, ECF subfamily   | Transcriptional regulators   |
| PA1365           | probable siderophore receptor   | Transport of small molecules   |
| PA1431           | regulatory protein RsaL   | Secreted Factors (toxins, enzymes, alginate)                               |
| A1556            | probable cytochrome c oxidase subunit   | Energy metabolism  |
| PA1559           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA1654           | probable aminotransferase   | Putative enzymes   |
| PA1655           | probable glutathione S-transferase  | Putative enzymes   |
| PA1687           | spermidine synthase   | Amino acid biosynthesis and metabolism                                     |
| PA1908           | probable major facilitator superfamily (MFS) transporter  | Membrane proteins  |
| PA1911           | probable transmembrane sensor   | Membrane proteins  |
| PA1917           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA1972           | conserved hypothetical protein  | Membrane proteins  |
| PA2002           | conserved hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA2021           | hypothetical protein  | Hypothetical, unclassified, unknown  |

| Product name  | Function class   |
|---|--|
| hypothetical protein  | Hypothetical, unclassified, unknown  |
| hypothetical protein  | Hypothetical, unclassified, unknown  |
| hypothetical protein  | Membrane proteins  |
| probable amino acid permease  | Transport of small molecules   |
| fimbrial subunit CupA4  | Motility & Attachment  |
| *   | Hypothetical, unclassified, unknown  |
|   | Secreted Factors (toxins, enzymes, alginate)   |
| · -   | Transcriptional regulators   |
|   | Hypothetical, unclassified, unknown  |
|   | Transcriptional regulators   |
|   | Membrane proteins  |
|   | Membrane proteins  |
|   | Membrane proteins  |
|   | Adaptation, Protection   |
|   | Hypothetical, unclassified, unknown  |
|   | Membrane proteins  |
|   | Hypothetical, unclassified, unknown  |
|   |  |
|   | Membrane proteins  |
|   | Hypothetical, unclassified, unknown  |
|   | Hypothetical, unclassified, unknown  |
| -   | Membrane proteins  |
|   | Hypothetical, unclassified, unknown  |
|   | Amino acid biosynthesis and metabolism   |
|   | Energy metabolism  |
| NADH dehydrogenase I chain B  | Energy metabolism  |
| two-component response regulator PfeR                               | Transcriptional regulators   |
| Ferric enterobactin receptor, outer membrane protein PfeA precursor | Membrane proteins  |
| conserved hypothetical protein                                      | Hypothetical, unclassified, unknown  |
| hypothetical protein  | Hypothetical, unclassified, unknown  |
| hypothetical protein  | Membrane proteins  |
| probable enoyl-CoA hydratase/isomerase                              | Putative enzymes   |
| conserved hypothetical protein                                      | Hypothetical, unclassified, unknown  |
| conserved hypothetical protein                                      | Hypothetical, unclassified, unknown  |
| hypothetical protein  | Membrane proteins  |
|   | Transcriptional regulators   |
| ribosome modulation factor  | Translation, post-translational modification, degradation  |
| histidinol-phosphate aminotransferase                               | Amino acid biosynthesis and metabolism   |
|   | Protein secretion/export apparatus   |
| -   | Membrane proteins  |
|   | Hypothetical, unclassified, unknown  |
|   | Membrane proteins  |
|   | Biosynthesis of cofactors, prosthetic groups and carriers  |
|   | Transport of small molecules   |
|   | Transport of small molecules   |
|   | Membrane proteins  |
|   | Transcriptional regulators   |
|   | · -  |
|   | Hypothetical, unclassified, unknown  |
|   | Hypothetical, unclassified, unknown  |
|   | Adaptation, Protection   |
|   | Adaptation, Protection   |
|   | Membrane proteins  |
|   | Membrane proteins  |
|   | Membrane proteins  |
|   | Hypothetical, unclassified, unknown  |
| •   | Putative enzymes   |
|   | Membrane proteins  |
| -   | Chaperones & heat shock proteins   |
| probable iron-binding protein IscA                                  | Biosynthesis of cofactors, prosthetic groups and carriers  |
| L-cysteine desulfurase (pyridoxal phosphate-dependent)              | Amino acid biosynthesis and metabolism   |
| hypothetical protein  | Hypothetical, unclassified, unknown  |
| Fe(III) dicitrate transport protein FecA                            | Membrane proteins  |
| hypothetical protein  | Hypothetical, unclassified, unknown  |
| 4-hydroxyphenylacetate 3-monooxygenase large chain                  | Carbon compound catabolism   |
|   |  |
| ferric enterobactin transport protein FepC                          | Transport of small molecules   |
| ferric enterobactin transport protein FepC<br>protease IV           | Secreted Factors (toxins, enzymes, alginate)   |
| protease IV   | Secreted Factors (toxins, enzymes, alginate)   |
|   |  |
|   | hypothetical protein<br>hypothetical protein<br>probable amino acid permease<br>fimbrial subunit CupA4<br>hypothetical protein<br>transcriptional regulator PtxR<br>probable transcriptional regulator<br>hypothetical protein<br>probable gima-70 factor, ECF subfamily<br>probable transmembrane sensor<br>probable ATP-binding/permease fusion ABC transporter<br>probable auimotrane protein precursor<br>Probable auimotrane protein precursor<br>PvdO<br>hypothetical protein<br>hypothetical protein<br>probable aminotransferase<br>NADH dehydrogenase I chain A<br>NADH dehydrogenase I chain B<br>two-component response regulator PfeR<br>Ferric enterobactin receptor, outer membrane protein PfeA precursor<br>conserved hypothetical protein<br>hypothetical protein<br>conserved hypothetical protein<br>hypothetical protein<br>ferredoxinNADP+ reductase<br>heme acquisition protein HasAp<br>Haem uptake outer membrane receptor HasR precursor<br>probable transmerbrane sensor<br>probable transmerbrane sensor<br>probable transmerbrane sensor<br>probable transmerbrane sensor<br>probable transmerbrane sensor<br>probable farior facilitator superfamily (MFS) transporter<br>conserved hypothetical protein<br>heat shock protein HasAB<br>probable major facilitator superfamily (MFS) tran |

| ORF              | Product name  | Function class  |  |  |
|------------------|---|---|--|--|
| PA4226           | dihydroaeruginoic acid synthetase                                   | Secreted Factors (toxins, enzymes, alginate)              |  |  |
| PA4227           | transcriptional regulator PchR                                      | Transcriptional regulators                                |  |  |
| PA4228           | pyochelin biosynthesis protein PchD                                 | Secreted Factors (toxins, enzymes, alginate)              |  |  |
| PA4229           | pyochelin biosynthetic protein PchC                                 | Secreted Factors (toxins, enzymes, alginate)              |  |  |
| PA4230           | salicylate biosynthesis protein PchB                                | Secreted Factors (toxins, enzymes, alginate)              |  |  |
| PA4231           | salicylate biosynthesis protein reinb                               | Secreted Factors (toxins, enzymes, alginate)              |  |  |
| PA4288           | probable transcriptional regulator                                  | Transcriptional regulators                                |  |  |
| PA4288<br>PA4357 | · · · ·   |   |  |  |
|                  | conserved hypothetical protein                                      | Hypothetical, unclassified, unknown                       |  |  |
| PA4358           | probable ferrous iron transport protein                             | Membrane proteins   |  |  |
| PA4359           | conserved hypothetical protein                                      | Hypothetical, unclassified, unknown                       |  |  |
| PA4370           | Insulin-cleaving metalloproteinase outer membrane protein precursor | Membrane proteins   |  |  |
| PA4371           | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |  |
| PA4372           | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |  |
| PA4500           | probable binding protein component of ABC transporter               | Transport of small molecules                              |  |  |
| PA4503           | probable permease of ABC transporter                                | Membrane proteins   |  |  |
| PA4513           | probable oxidoreductase   | Putative enzymes  |  |  |
| PA4514           | probable outer membrane receptor for iron transport                 | Transport of small molecules                              |  |  |
| PA4515           | conserved hypothetical protein                                      | Hypothetical, unclassified, unknown                       |  |  |
| PA4516           | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |  |
| PA4517           | conserved hypothetical protein                                      | Hypothetical, unclassified, unknown                       |  |  |
| PA4519           | ornithine decarboxylase   | Amino acid biosynthesis and metabolism                    |  |  |
| PA4629           | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |  |
| PA4630           | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |  |
| PA4633           | probable chemotaxis transducer                                      | Adaptation, Protection                                    |  |  |
| PA4654           | probable major facilitator superfamily (MFS) transporter            | Membrane proteins   |  |  |
| PA4675           | probable TonB-dependent receptor                                    | Transport of small molecules                              |  |  |
| PA4687           | ferric iron-binding periplasmic protein HitA                        | Transport of small molecules                              |  |  |
|                  |   |   |  |  |
| PA4688           | iron (III)-transport system permease HitB                           | Membrane proteins   |  |  |
| PA4706           | probable ATP-binding component of ABC transporter                   | Transport of small molecules                              |  |  |
| PA4773           | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |  |
| PA4785           | probable acyl-CoA thiolase  | Putative enzymes  |  |  |
| PA4826           | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |  |
| PA4833           | conserved hypothetical protein                                      | Membrane proteins   |  |  |
| PA4838           | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |  |
| PA4858           | conserved hypothetical protein                                      | Hypothetical, unclassified, unknown                       |  |  |
| PA4860           | probable permease of ABC transporter                                | Membrane proteins   |  |  |
| PA4861           | probable ATP-binding component of ABC transporter                   | Transport of small molecules                              |  |  |
| PA4862           | probable ATP-binding component of ABC transporter                   | Transport of small molecules                              |  |  |
| PA4883           | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |  |
| PA4892           | urease accessory protein UreF                                       | Biosynthesis of cofactors, prosthetic groups and carriers |  |  |
| PA4893           | urease accessory protein UreG                                       | Biosynthesis of cofactors, prosthetic groups and carriers |  |  |
| PA4894           | hypothetical protein  | Membrane proteins   |  |  |
| PA4895           | probable transmembrane sensor                                       | Membrane proteins   |  |  |
| PA4897           | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |  |
| PA4973           | thiamin biosynthesis protein ThiC                                   | Biosynthesis of cofactors, prosthetic groups and carriers |  |  |
| PA5119           | glutamine synthetase  | Amino acid biosynthesis and metabolism                    |  |  |
| PA5124           | two-component sensor NtrB   | Two-component regulatory systems                          |  |  |
| PA5125           | two-component response regulator NtrC                               | Transcriptional regulators                                |  |  |
| PA5150           | probable short-chain dehydrogenase                                  | Putative enzymes  |  |  |
| PA5217           | probable binding protein component of ABC iron transporter          | Transport of small molecules                              |  |  |
| PA5287           | ammonium transporter AmtB   | Membrane proteins   |  |  |
| PA5287<br>PA5309 | probable oxidoreductase   | Putative enzymes  |  |  |
|                  | 1   | -   |  |  |
| PA5312           | probable aldehyde dehydrogenase                                     | Putative enzymes  |  |  |
| PA5313           | probable pyridoxal-dependent aminotransferase                       | Putative enzymes  |  |  |
| PA5314           | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |  |
| PA5386           | probable 3-hydroxyacyl-CoA dehydrogenase                            | Putative enzymes  |  |  |
| PA5479           | proton-glutamate symporter  | Membrane proteins   |  |  |
| PA5526           | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |  |
| PA5531           | TonB protein  | Transport of small molecules                              |  |  |

#### **Quorum Sensing**

| PA0006           | conserved hypothetical protein  |
|------------------|---|
| PA0048           | probable transcriptional regulator  |
| PA0049           | hypothetical protein  |
| PA0050<br>PA0051 | hypothetical protein  |
| PA0051<br>PA0052 | potential phenazine-modifying enzyme<br>hypothetical protein                |
| PA0052<br>PA0055 | hypothetical protein  |
| PA0079           | hypothetical protein  |
| PA0080           | hypothetical protein  |
| PA0097           | hypothetical protein  |
| PA0105           | cytochrome c oxidase, subunit II  |
| PA0106           | cytochrome c oxidase, subunit I   |
| PA0107           | conserved hypothetical protein  |
| PA0108           | cytochrome c oxidase, subunit III   |
| PA0112           | hypothetical protein  |
| PA0113           | probable cytochrome c oxidase assembly factor                               |
| PA0119           | probable dicarboxylate transporter  |
| PA0122<br>PA0129 | conserved hypothetical protein<br>gamma-aminobutyrate permease              |
| PA0129<br>PA0130 | probable aldehyde dehydrogenase   |
| PA0130           | hypothetical protein  |
| PA0132           | beta-alaninepyruvate transaminase   |
| PA0144           | hypothetical protein  |
| PA0145           | hypothetical protein  |
| PA0153           | protocatechuate 3,4-dioxygenase, beta subunit                               |
| PA0154           | protocatechuate 3,4-dioxygenase, alpha subunit                              |
| PA0160           | hypothetical protein  |
| PA0167           | probable transcriptional regulator  |
| PA0191           | probable transcriptional regulator  |
| PA0200           | hypothetical protein  |
| PA0209           | conserved hypothetical protein  |
| PA0210<br>PA0211 | malonate decarboxylase delta subunit<br>malonate decarboxylase beta subunit |
| PA0211<br>PA0212 | malonate decarboxylase gamma subunit  |
| PA0212           | hypothetical protein  |
| PA0214           | probable acyl transferase   |
| PA0221           | probable aminotransferase   |
| PA0226           | probable CoA transferase, subunit A   |
| PA0227           | probable CoA transferase, subunit B   |
| PA0228           | beta-ketoadipyl CoA thiolase PcaF   |
| PA0234           | hypothetical protein  |
| PA0235           | 4-hydroxybenzoate transporter PcaK  |
| PA0239           | hypothetical protein  |
| PA0241           | probable major facilitator superfamily (MFS) transporter                    |
| PA0251<br>PA0264 | hypothetical protein<br>hypothetical protein                                |
| PA0265           | succinate-semialdehyde dehydrogenase  |
| PA0266           | 4-aminobutyrate aminotransferase  |
| PA0269           | conserved hypothetical protein  |
| PA0270           | hypothetical protein  |
| PA0271           | hypothetical protein  |
| PA0274           | hypothetical protein  |
| PA0287           | 3-guanidinopropionate transport protein                                     |
| PA0297           | probable glutamine amidotransferase   |
| PA0298           | probable glutamine synthetase   |
| PA0299           | putrescine aminotransferase   |
| PA0300           | polyamine transport protein   |
| PA0301<br>PA0302 | polyamine transport protein<br>polyamine transport protein PotG             |
| PA0302<br>PA0303 | polyamine transport protein PotG  |
| PA0303<br>PA0304 | polyamine transport protein Polifi<br>polyamine transport protein Polifi    |
| PA0304<br>PA0305 | hypothetical protein  |
| PA0321           | probable acetylpolyamine aminohydrolase                                     |
| PA0363           | phosphopantetheine adenylyltransferase                                      |
| PA0364           | probable oxidoreductase   |
| PA0385           | hypothetical protein  |
|                  |   |

#### Function class

Hypothetical, unclassified, unknown Transcriptional regulators Hypothetical, unclassified, unknown Hypothetical, unclassified, unknown Putative enzymes Hypothetical, unclassified, unknown Energy metabolism Energy metabolism Energy metabolism Energy metabolism Membrane proteins Energy metabolism Transport of small molecules Hypothetical, unclassified, unknown Transport of small molecules Putative enzymes Hypothetical, unclassified, unknown Amino acid biosynthesis and metabolism Hypothetical, unclassified, unknown Hypothetical, unclassified, unknown Carbon compound catabolism Carbon compound catabolism Hypothetical, unclassified, unknown Transcriptional regulators Transcriptional regulators Hypothetical, unclassified, unknown Putative enzymes Carbon compound catabolism Putative enzymes Carbon compound catabolism Carbon compound catabolism Fatty acid and phospholipid metabolism Hypothetical, unclassified, unknown Membrane proteins Membrane proteins Membrane proteins Hypothetical, unclassified, unknown Hypothetical, unclassified, unknown Central intermediary metabolism Central intermediary metabolism Hypothetical, unclassified, unknown Hypothetical, unclassified, unknown Hypothetical, unclassified, unknown Hypothetical, unclassified, unknown Transport of small molecules Amino acid biosynthesis and metabolism Putative enzymes Putative enzymes Transport of small molecules Transport of small molecules Transport of small molecules Membrane proteins Membrane proteins Hypothetical, unclassified, unknown Putative enzymes Central intermediary metabolism Putative enzymes Hypothetical, unclassified, unknown

| ORF              | Product name                                     | Function class  |
|------------------|--|---|
| PA0399           | cystathionine beta-synthase                      | Amino acid biosynthesis and metabolism                    |
| PA0436           | probable transcriptional regulator               | Transcriptional regulators                                |
| PA0439           | probable oxidoreductase                          | Putative enzymes  |
| PA0440           | probable oxidoreductase                          | Putative enzymes  |
| PA0441           | dihydropyrimidinase                              | Nucleotide biosynthesis and metabolism                    |
| PA0443           | probable transporter                             | Membrane proteins   |
| PA0444           | N-carbamoyl-beta-alanine amidohydrolase          | Nucleotide biosynthesis and metabolism                    |
| PA0453           | hypothetical protein                             | Hypothetical, unclassified, unknown                       |
| PA0456           | probable cold-shock protein                      | Transcriptional regulators                                |
| PA0459           | probable ClpA/B protease ATP binding subunit     | Translation, post-translational modification, degradation |
| PA0467           | conserved hypothetical protein                   | Hypothetical, unclassified, unknown                       |
| PA0468           | hypothetical protein                             | Hypothetical, unclassified, unknown                       |
| PA0468<br>PA0469 | hypothetical protein                             | Hypothetical, unclassified, unknown                       |
|                  | 51 1   |   |
| PA0473           | probable glutathione S-transferase               | Putative enzymes  |
| PA0474           | hypothetical protein                             | Hypothetical, unclassified, unknown                       |
| PA0480           | probable hydrolase                               | Putative enzymes  |
| PA0482           | malate synthase G                                | Central intermediary metabolism                           |
| PA0489           | probable phosphoribosyl transferase              | Putative enzymes  |
| PA0497           | hypothetical protein                             | Hypothetical, unclassified, unknown                       |
| PA0498           | hypothetical protein                             | Hypothetical, unclassified, unknown                       |
| PA0512           | conserved hypothetical protein                   | Biosynthesis of cofactors, prosthetic groups and carriers |
| PA0513           | probable transcriptional regulator               | Biosynthesis of cofactors, prosthetic groups and carriers |
| PA0527           | transcriptional regulator Dnr                    | Transcriptional regulators                                |
| PA0542           | conserved hypothetical protein                   | Hypothetical, unclassified, unknown                       |
| PA0610           | transcriptional regulator PrtN                   | Transcriptional regulators                                |
| PA0614           | hypothetical protein                             | Hypothetical, unclassified, unknown                       |
| PA0615           | hypothetical protein                             | Hypothetical, unclassified, unknown                       |
| PA0616           | hypothetical protein                             | Related to phage, transposon, or plasmid                  |
| PA0617           | probable bacteriophage protein                   | Related to phage, transposon, or plasmid                  |
| PA0618           | probable bacteriophage protein                   | Related to phage, transposon, or plasmid                  |
| PA0619           | probable bacteriophage protein                   | Related to phage, transposon, or plasmid                  |
| PA0620           | probable bacteriophage protein                   | Related to phage, transposon, or plasmid                  |
| PA0621           | conserved hypothetical protein                   | Related to phage, transposon, or plasmid                  |
|                  |  |   |
| PA0622           | probable bacteriophage protein                   | Related to phage, transposon, or plasmid                  |
| PA0623           | probable bacteriophage protein                   | Related to phage, transposon, or plasmid                  |
| PA0624           | hypothetical protein                             | Related to phage, transposon, or plasmid                  |
| PA0625           | hypothetical protein                             | Related to phage, transposon, or plasmid                  |
| PA0626           | hypothetical protein                             | Related to phage, transposon, or plasmid                  |
| PA0627           | conserved hypothetical protein                   | Related to phage, transposon, or plasmid                  |
| PA0628           | conserved hypothetical protein                   | Related to phage, transposon, or plasmid                  |
| PA0629           | conserved hypothetical protein                   | Related to phage, transposon, or plasmid                  |
| PA0630           | hypothetical protein                             | Related to phage, transposon, or plasmid                  |
| PA0631           | hypothetical protein                             | Related to phage, transposon, or plasmid                  |
| PA0632           | hypothetical protein                             | Related to phage, transposon, or plasmid                  |
| PA0635           | hypothetical protein                             | Related to phage, transposon, or plasmid                  |
| PA0636           | hypothetical protein                             | Related to phage, transposon, or plasmid                  |
| PA0637           | conserved hypothetical protein                   | Related to phage, transposon, or plasmid                  |
| PA0638           | probable bacteriophage protein                   | Related to phage, transposon, or plasmid                  |
| PA0639           | conserved hypothetical protein                   | Related to phage, transposon, or plasmid                  |
| PA0640           | probable bacteriophage protein                   | Related to phage, transposon, or plasmid                  |
| PA0641           | probable bacteriophage protein                   | Related to phage, transposon, or plasmid                  |
| PA0642           | hypothetical protein                             | Related to phage, transposon, or plasmid                  |
| PA0643           | hypothetical protein                             | Related to phage, transposon, or plasmid                  |
| PA0644           | hypothetical protein                             | Related to phage, transposon, or plasmid                  |
| PA0645           | hypothetical protein                             | Related to phage, transposon, or plasmid                  |
| PA0646           | hypothetical protein                             | Related to phage, transposon, or plasmid                  |
| PA0647           | hypothetical protein                             | Related to phage, transposon, or plasmid                  |
| PA0648           | hypothetical protein                             | Related to phage, transposon, or plasmid                  |
| PA0671           | hypothetical protein                             | Hypothetical, unclassified, unknown                       |
|                  | probable type II secretion system protein        | Protein secretion/export apparatus                        |
| PA0683           |  |   |
| PA0701           | probable transcriptional regulator               | Transcriptional regulators                                |
| PA0717           | hypothetical protein of bacteriophage Pf1        | Hypothetical, unclassified, unknown                       |
| PA0718           | hypothetical protein of bacteriophage Pf1        | Hypothetical, unclassified, unknown                       |
| PA0719           | hypothetical protein of bacteriophage Pf1        | Hypothetical, unclassified, unknown                       |
| PA0720           | helix destabilizing protein of bacteriophage Pf1 | DNA replication, recombination, modification and repair   |
| PA0721           | hypothetical protein of bacteriophage Pf1        | Hypothetical, unclassified, unknown                       |
| PA0723           | coat protein B of bacteriophage Pf1              | Related to phage, transposon, or plasmid                  |
| PA0736           | hypothetical protein                             | Membrane proteins   |

| ORF              | Product name  | Function class  |
|------------------|---|---|
| PA0740           | SDS hydrolase SdsA1   | Putative enzymes  |
| PA0745           | probable enoyl-CoA hydratase/isomerase                          | Putative enzymes  |
| PA0746           | probable acyl-CoA dehydrogenase                                 | Putative enzymes  |
| PA0747           | probable aldehyde dehydrogenase                                 | Putative enzymes  |
| PA0776           | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA0783           | sodium/proline symporter PutP                                   | Membrane proteins   |
| PA0826           | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA0841           | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA0845           | conserved hypothetical protein                                  | Hypothetical, unclassified, unknown                       |
| PA0854           | fumarate hydratase  | Energy metabolism   |
| PA0867           | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA0870           | aromatic amino acid aminotransferase                            | Amino acid biosynthesis and metabolism                    |
| PA0871           | pterin-4-alpha-carbinolamine dehydratase                        | Amino acid biosynthesis and metabolism                    |
| PA0885           | probable C4-dicarboxylate transporter                           | Transport of small molecules                              |
| PA0910           | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA0938           | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA0943           | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA0997           | Homologous to beta-keto-acyl-acyl-carrier protein synthase      | Biosynthesis of cofactors, prosthetic groups and carriers |
| PA0997           | Homologous to beta-keto-acyl-acyl-carrier protein synthase      | Biosynthesis of cofactors, prosthetic groups and carriers |
| PA0998           | 3-oxoacyl-[acyl-carrier-protein] synthase III                   | Biosynthesis of cofactors, prosthetic groups and carriers |
|                  | hypothetical protein  |   |
| PA1018           | 51 1  | Hypothetical, unclassified, unknown                       |
| PA1019           | cis,cis-muconate transporter MucK                               | Membrane proteins   |
| PA1025           | probable porin  | Membrane proteins   |
| PA1049           | pyridoxine 5'-phosphate oxidase                                 | Biosynthesis of cofactors, prosthetic groups and carriers |
| PA1051           | probable transporter  | Membrane proteins   |
| PA1070           | branched-chain amino acid transport protein BraG                | Transport of small molecules                              |
| PA1072           | branched-chain amino acid transport protein BraE                | Membrane proteins   |
| PA1073           | branched-chain amino acid transport protein BraD                | Membrane proteins   |
| PA1074           | branched-chain amino acid transport protein BraC                | Transport of small molecules                              |
| PA1106           | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA1113           | probable ATP-binding/permease fusion ABC transporter            | Membrane proteins   |
| PA1123           | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA1127           | probable oxidoreductase   | Adaptation, Protection                                    |
| PA1130           | rhamnosyltransferase 2  | Cell wall / LPS / capsule                                 |
| PA1131           | probable major facilitator superfamily (MFS) transporter        | Membrane proteins   |
| PA1137           | probable oxidoreductase   | Putative enzymes  |
| PA1143           | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA1144           | probable major facilitator superfamily (MFS) transporter        | Membrane proteins   |
| PA1150           | pyocin S2   | Adaptation, Protection                                    |
| PA1166           | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA1183           | C4-dicarboxylate transport protein                              | Transport of small molecules                              |
| PA1186           | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA1190           | conserved hypothetical protein                                  | Membrane proteins   |
| PA1197           | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA1206           | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA1207           | glutathione-regulated potassium-efflux system protein KefB      | Transport of small molecules                              |
| PA1220           | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA1223           | probable transcriptional regulator                              | Transcriptional regulators                                |
| PA1228           | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA1236           | probable major facilitator superfamily (MFS) transporter        | Membrane proteins   |
| PA1239           | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA1242           | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA1250           | alkaline proteinase inhibitor AprI                              | Secreted Factors (toxins, enzymes, alginate)              |
| PA1262           | probable major facilitator superfamily (MFS) transporter        | Membrane proteins   |
| PA1266           | probable oxidoreductase   | Putative enzymes  |
| PA1338           | gamma-glutamyltranspeptidase precursor                          | Amino acid biosynthesis and metabolism                    |
| PA1340           | probable permease of ABC transporter                            | Transport of small molecules                              |
| PA1341           | probable permease of ABC transporter                            | Membrane proteins   |
| PA1342           | probable binding protein component of ABC transporter           | Transport of small molecules                              |
| PA1344           | probable short-chain dehydrogenase                              | Putative enzymes  |
| PA1367           | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA1377           | conserved hypothetical protein                                  | Hypothetical, unclassified, unknown                       |
| PA1379           | probable short-chain dehydrogenase                              | Putative enzymes  |
| PA1386           | probable ATP-binding component of ABC transporter               | Transport of small molecules                              |
| PA1395           | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA1393<br>PA1399 | probable transcriptional regulator                              | Transcriptional regulators                                |
|                  |   | Energy metabolism   |
| DA 1/100         |   | LUEISV IIIEIADUUNII                                       |
| PA1400<br>PA1416 | probable pyruvate carboxylase<br>conserved hypothetical protein | Hypothetical, unclassified, unknown                       |

| ORF              | Product name  | Function class   |
|------------------|---|--|
| PA1418           | probable sodium:solute symport protein  | Transport of small molecules   |
| PA1419           | probable transporter  | Membrane proteins  |
| PA1426           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA1431           | regulatory protein RsaL   | Secreted Factors (toxins, enzymes, alginate)                               |
| PA1440           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA1519           | probable transporter  | Membrane proteins  |
| PA1537           | probable short-chain dehydrogenase  | Putative enzymes   |
| PA1538           | probable flavin-containing monooxygenase  | Putative enzymes   |
| PA1540           | conserved hypothetical protein  | Membrane proteins  |
| PA1542           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA1568           | conserved hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA1569           | probable major facilitator superfamily (MFS) transporter                        | Membrane proteins  |
| PA1575           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA1576           | probable 3-hydroxyisobutyrate dehydrogenase                                     | Putative enzymes   |
| PA1608           | probable chemotaxis transducer  | Adaptation, Protection   |
| PA1609           | beta-ketoacyl-ACP synthase I  | Fatty acid and phospholipid metabolism                                     |
| PA1610           | beta-hydroxydecanoyl-ACP dehydrase  | Fatty acid and phospholipid metabolism                                     |
| PA1617           | probable AMP-binding enzyme   | Putative enzymes   |
| PA1650           | probable transporter  | Membrane proteins  |
| PA1746           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA1759           | probable transcriptional regulator  | Transcriptional regulators   |
| PA1763           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA1768           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA1772           | probable methyltransferase  | Biosynthesis of cofactors, prosthetic groups and carriers                  |
| PA1774           | CfrX protein  | Hypothetical, unclassified, unknown  |
| PA1775           | conserved cytoplasmic membrane protein, CmpX protein                            | Membrane proteins  |
| PA1776           | ECF sigma factor SigX   | Transcriptional regulators   |
| PA1781           | assimilatory nitrite reductase large subunit                                    | Central intermediary metabolism  |
| PA1784<br>PA1789 | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA1789           | hypothetical protein<br>hypothetical protein                                    | Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown |
| PA1790<br>PA1812 | membrane-bound lytic murein transglycosylase D precursor                        | Amino acid biosynthesis and metabolism                                     |
| PA1812<br>PA1824 | conserved hypothetical protein  | Membrane proteins  |
| PA1852           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA1853           | probable transcriptional regulator  | Transcriptional regulators   |
| PA1871           | LasA protease precursor   | Secreted Factors (toxins, enzymes, alginate)                               |
| PA1875           | probable outer membrane protein precursor                                       | Protein secretion/export apparatus   |
| PA1891           | hypothetical protein  | Membrane proteins  |
| PA1892           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA1893           | hypothetical protein  | Putative enzymes   |
| PA1895           | hypothetical protein  | Membrane proteins  |
| PA1896           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA1897           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA1898           | quorum-sensing control repressor  | Transcriptional regulators   |
| PA1901           | phenazine biosynthesis protein PhzC   | Secreted Factors (toxins, enzymes, alginate)                               |
| PA1903           | phenazine biosynthesis protein PhzE   | Secreted Factors (toxins, enzymes, alginate)                               |
| PA1905           | probable pyridoxamine 5'-phosphate oxidase                                      | Secreted Factors (toxins, enzymes, alginate)                               |
| PA1922           | probable TonB-dependent receptor  | Transport of small molecules   |
| PA1924           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA1927           | 5-methyltetrahydropteroyltriglutamate-homocysteine S-methyltransferase          | Amino acid biosynthesis and metabolism                                     |
| PA1935           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA1942           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA1946           | binding protein component precursor of ABC ribose transporter                   | Transport of small molecules   |
| PA1947           | ribose transport protein RbsA   | Transport of small molecules   |
| PA1955           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA1974           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA1976           | probable two-component sensor   | Two-component regulatory systems   |
| PA1977           | hypothetical protein  | Membrane proteins  |
| PA1978           | probable transcriptional regulator  | Transcriptional regulators   |
| PA1979           | probable two-component sensor   | Two-component regulatory systems   |
| PA1980           | probable two-component response regulator                                       | Transcriptional regulators   |
| PA1981           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA1983<br>PA1984 | cytochrome c550   | Energy metabolism  |
| PATYXA           | probable aldehyde dehydrogenase   | Putative enzymes   |
|                  |   | Destation and an an  |
| PA1991           | probable iron-containing alcohol dehydrogenase                                  | Putative enzymes   |
| PA1991<br>PA1992 | probable iron-containing alcohol dehydrogenase<br>probable two-component sensor | Two-component regulatory systems   |
| PA1991           | probable iron-containing alcohol dehydrogenase                                  |  |

| ORF              | Product name  | Function class   |
|------------------|---|--|
| PA2002           | conserved hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA2003           | 3-hydroxybutyrate dehydrogenase   | Carbon compound catabolism   |
| PA2004           | conserved hypothetical protein  | Membrane proteins  |
| PA2006           | probable major facilitator superfamily (MFS) transporter                                  | Membrane proteins  |
| PA2012           | methylcrotonyl-CoA carboxylase, alpha-subunit (biotin-containing)                         | Carbon compound catabolism   |
| PA2013<br>PA2014 | putative 3-methylglutaconyl-CoA hydratase<br>methylcrotonyl-CoA carboxylase, beta-subunit | Carbon compound catabolism<br>Carbon compound catabolism                   |
| PA2014           | putative isovaleryl-CoA dehydrogenase   | Carbon compound catabolism   |
| PA2018           | Resistance-Nodulation-Cell Division (RND) multidrug efflux transporter                    | Transport of small molecules   |
| PA2030           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA2031           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA2041           | probable amino acid permease  | Transport of small molecules   |
| PA2060           | probable permease of ABC transporter  | Transport of small molecules   |
| PA2066<br>PA2067 | hypothetical protein<br>probable hydrolase  | Hypothetical, unclassified, unknown<br>Putative enzymes                    |
| PA2067<br>PA2068 | probable major facilitator superfamily (MFS) transporter                                  | Membrane proteins  |
| PA2069           | probable carbamoyl transferase  | Putative enzymes   |
| PA2079           | probable amino acid permease  | Transport of small molecules   |
| PA2091           | hypothetical protein  | Membrane proteins  |
| PA2109           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA2110           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA2116<br>PA2119 | conserved hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA2119<br>PA2122 | alcohol dehydrogenase (Zn-dependent)<br>hypothetical protein                              | Putative enzymes<br>Hypothetical, unclassified, unknown                    |
| PA2129           | chaperone CupA2   | Motility & Attachment  |
| PA2134           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA2160           | probable glycosyl hydrolase   | Putative enzymes   |
| PA2161           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA2169           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA2172           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA2185           | non-heme catalase KatN  | Adaptation, Protection   |
| PA2198<br>PA2201 | hypothetical protein<br>hypothetical protein  | Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown |
| PA2209           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA2210           | probable major facilitator superfamily (MFS) transporter                                  | Membrane proteins  |
| PA2211           | conserved hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA2212           | conserved hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA2213           | probable porin  | Membrane proteins  |
| PA2214           | probable major facilitator superfamily (MFS) transporter                                  | Membrane proteins  |
| PA2217<br>PA2225 | probable aldehyde dehydrogenase<br>hypothetical protein                                   | Putative enzymes<br>Hypothetical, unclassified, unknown                    |
| PA2228           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA2249           | branched-chain alpha-keto acid dehydrogenase (lipoamide component)                        | Amino acid biosynthesis and metabolism                                     |
| PA2250           | lipoamide dehydrogenase-Val   | Amino acid biosynthesis and metabolism                                     |
| PA2258           | transcriptional regulator PtxR  | Secreted Factors (toxins, enzymes, alginate)                               |
| PA2274           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA2300           | chitinase   | Carbon compound catabolism   |
| PA2303<br>PA2313 | hypothetical protein<br>hypothetical protein  | Hypothetical, unclassified, unknown<br>Membrane proteins                   |
| PA2313<br>PA2322 | gluconate permease  | Transport of small molecules   |
| PA2328           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA2329           | probable ATP-binding component of ABC transporter   | Transport of small molecules   |
| PA2330           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA2334           | probable transcriptional regulator  | Transcriptional regulators   |
| PA2359           | probable transcriptional regulator  | Transcriptional regulators   |
| PA2423           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA2439<br>PA2453 | hypothetical protein<br>hypothetical protein  | Membrane proteins<br>Hypothetical, unclassified, unknown                   |
| PA2455<br>PA2474 | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA2499           | probable deaminase  | Putative enzymes   |
| PA2507           | catechol 1,2-dioxygenase  | Carbon compound catabolism   |
| PA2508           | muconolactone delta-isomerase   | Carbon compound catabolism   |
| PA2509           | muconate cycloisomerase I   | Carbon compound catabolism   |
| PA2511           | probable transcriptional regulator  | Transcriptional regulators   |
| PA2512           | anthranilate dioxygenase large subunit  | Carbon compound catabolism   |
| PA2513<br>PA2514 | anthranilate dioxygenase small subunit<br>anthranilate dioxygenase reductase              | Carbon compound catabolism<br>Carbon compound catabolism                   |
| PA2514<br>PA2515 | cis-1,2-dihydroxycyclohexa-3,4-diene carboxylate dehydrogenase                            | Carbon compound catabolism   |
| 1114313          | and the anithmory of context of the carboxy and achydrogenase                             | Carbon compound cambonism  |

| ORF              | Product name   | Function class   |
|------------------|--|--|
| PA2516           | toluate 1,2-dioxygenase electron transfer component  | Energy metabolism  |
| PA2517           | toluate 1,2-dioxygenase beta subunit   | Carbon compound catabolism   |
| PA2518           | toluate 1,2-dioxygenase alpha subunit  | Carbon compound catabolism   |
| PA2519           | transcriptional regulator XylS   | Transcriptional regulators   |
| PA2529           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA2554           | probable short-chain dehydrogenase   | Putative enzymes   |
| PA2555           | probable AMP-binding enzyme  | Putative enzymes   |
| PA2557           | probable AMP-binding enzyme  | Fatty acid and phospholipid metabolism   |
| PA2564<br>PA2566 | hypothetical protein<br>conserved hypothetical protein   | Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown                       |
| PA2567           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA2570           | LecA   | Adaptation, Protection   |
| PA2587           | probable FAD-dependent monooxygenase   | Biosynthesis of cofactors, prosthetic groups and carriers  |
| PA2591           | probable transcriptional regulator   | Transcriptional regulators   |
| PA2595           | conserved hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA2625           | conserved hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA2629           | adenylosuccinate lyase   | Amino acid biosynthesis and metabolism   |
| PA2654           | probable chemotaxis transducer   | Adaptation, Protection   |
| PA2660           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA2667<br>PA2682 | conserved hypothetical protein   | Hypothetical, unclassified, unknown<br>Putative enzymes  |
| PA2682<br>PA2711 | conserved hypothetical protein<br>probable periplasmic spermidine/putrescine-binding protein           | Transport of small molecules   |
| PA2717           | chloroperoxidase precursor   | Central intermediary metabolism  |
| PA2719           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA2753           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA2754           | conserved hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA2769           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA2776           | conserved hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA2778           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA2779           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA2780<br>PA2781 | hypothetical protein<br>hypothetical protein   | Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown                       |
| PA2801           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA2805           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA2819           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA2838           | probable transcriptional regulator   | Transcriptional regulators   |
| PA2847           | conserved hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA2851           | translation elongation factor P  | Translation, post-translational modification, degradation  |
| PA2876           | orotidine 5'-phosphate decarboxylase   | Nucleotide biosynthesis and metabolism   |
| PA2892           | GCase, alpha-subunit (biotin-containing)   | Putative enzymes   |
| PA2905<br>PA2915 | precorrin isomerase CobH<br>hypothetical protein   | Biosynthesis of cofactors, prosthetic groups and carriers<br>Hypothetical, unclassified, unknown |
| PA2922           | probable hydrolase   | Putative enzymes   |
| PA2924           | histidine transport system permease HisQ   | Membrane proteins  |
| PA2927           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA2938           | probable transporter   | Membrane proteins  |
| PA2939           | probable aminopeptidase  | Secreted Factors (toxins, enzymes, alginate)   |
| PA2942           | probable magnesium chelatase   | Biosynthesis of cofactors, prosthetic groups and carriers  |
| PA2950           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA2960           | type 4 fimbrial biogenesis protein PilZ  | Motility & Attachment  |
| PA2966<br>PA2967 | acyl carrier protein<br>3-oxoacyl-[acyl-carrier-protein] reductase                                     | Fatty acid and phospholipid metabolism<br>Fatty acid and phospholipid metabolism                 |
| PA2967<br>PA2968 | malonyl-CoA-[acyl-carrier-protein] transacylase  | Fatty acid and phospholipid metabolism   |
| PA3032           | cytochrome c Snr1  | Energy metabolism  |
| PA3035           | probable glutathione S-transferase   | Putative enzymes   |
| PA3036           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA3062           | PelC   | Cell wall / LPS / capsule  |
| PA3066           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA3079           | hypothetical protein   | Membrane proteins  |
| PA3131           | probable aldolase  | Central intermediary metabolism  |
| PA3139           | probable amino acid aminotransferase   | Amino acid biosynthesis and metabolism   |
| PA3186<br>PA3188 | Glucose/carbohydrate outer membrane porin OprB precursor<br>probable permease of ABC sugar transporter | Transport of small molecules<br>Transport of small molecules                                     |
| PA3188           | probable permease of ABC sugar transporter   | Transport of small molecules   |
| PA3208           | conserved hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA3209           | conserved hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA3213           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| 1110210          |  |  |

| ORF              | Product name  | Function class  |
|------------------|---|---|
| PA3234           | probable sodium:solute symporter  | Membrane proteins   |
| PA3236           | probable glycine betaine-binding protein precursor                              | Adaptation, Protection  |
| PA3237           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA3249           | probable transcriptional regulator  | Transcriptional regulators  |
| PA3270           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA3283           | conserved hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA3295           | probable HIT family protein   | Putative enzymes  |
| PA3301           | hypothetical protein  | Putative enzymes  |
| PA3309           | conserved hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA3313<br>PA3326 | hypothetical protein<br>probable Clp-family ATP-dependent protease              | Transport of small molecules<br>Translation, post-translational modification, degradation |
| PA3329           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA3330           | probable short chain dehydrogenase  | Putative enzymes  |
| PA3331           | cytochrome P450   | Adaptation, Protection  |
| PA3332           | conserved hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA3333           | 3-oxoacyl-[acyl-carrier-protein] synthase III                                   | Fatty acid and phospholipid metabolism  |
| PA3334           | probable acyl carrier protein   | Fatty acid and phospholipid metabolism  |
| PA3335           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA3336           | probable major facilitator superfamily (MFS) transporter                        | Membrane proteins   |
| PA3337           | ADP-L-glycero-D-mannoheptose 6-epimerase  | Cell wall / LPS / capsule   |
| PA3361           | fucose-binding lectin PA-IIL  | Motility & Attachment   |
| PA3362           | hypothetical protein<br>aliphatic amidase regulator                             | Membrane proteins<br>Carbon compound catabolism   |
| PA3363<br>PA3364 | aliphatic amidase regulator<br>aliphatic amidase expression-regulating protein  | Transcriptional regulators  |
| PA3365           | probable chaperone  | Chaperones & heat shock proteins  |
| PA3366           | aliphatic amidase   | Carbon compound catabolism  |
| PA3394           | NosF protein  | Energy metabolism   |
| PA3422           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA3431           | conserved hypothetical protein  | Membrane proteins   |
| PA3432           | hypothetical protein  | Membrane proteins   |
| PA3458           | probable transcriptional regulator  | Transcriptional regulators  |
| PA3466           | probable ATP-dependent RNA helicase   | Transcription, RNA processing and degradation   |
| PA3467           | probable major facilitator superfamily (MFS) transporter                        | Membrane proteins   |
| PA3478           | rhamnosyltransferase chain B  | Secreted Factors (toxins, enzymes, alginate)  |
| PA3479<br>PA3480 | rhamnosyltransferase chain A<br>probable deoxycytidine triphosphate deaminase   | Secreted Factors (toxins, enzymes, alginate)<br>Nucleotide biosynthesis and metabolism    |
| PA3480<br>PA3507 | probable short-chain dehydrogenase  | Putative enzymes  |
| PA3524           | lactoylglutathione lyase  | Central intermediary metabolism   |
| PA3529           | probable peroxidase   | Adaptation, Protection  |
| PA3535           | probable serine protease  | Putative enzymes  |
| PA3542           | alginate biosynthesis protein Alg44   | Cell wall / LPS / capsule   |
| PA3555           | AmD   | Adaptation, Protection  |
| PA3559           | probable nucleotide sugar dehydrogenase   | Putative enzymes  |
| PA3560           | phosphotransferase system, fructose-specific IIBC component                     | Carbon compound catabolism  |
| PA3562           | probable phosphotransferase system enzyme I                                     | Central intermediary metabolism   |
| PA3564           | conserved hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA3565<br>PA3568 | probable transcriptional regulator  | Transcriptional regulators  |
| PA3568<br>PA3569 | probable acetyl-coa synthetase<br>3-hydroxyisobutyrate dehydrogenase            | Putative enzymes<br>Carbon compound catabolism  |
| PA3509<br>PA3570 | methylmalonate-semialdehyde dehydrogenase                                       | Amino acid biosynthesis and metabolism  |
| PA3572           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA3581           | glycerol uptake facilitator protein   | Transport of small molecules  |
| PA3582           | glycerol kinase   | Central intermediary metabolism   |
| PA3586           | probable hydrolase  | Putative enzymes  |
| PA3595           | probable major facilitator superfamily (MFS) transporter                        | Membrane proteins   |
| PA3608           | polyamine transport protein PotB  | Membrane proteins   |
| PA3613           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA3614           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA3618           | conserved hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA3621<br>PA3661 | ferredoxin I<br>hypothetical protein  | Energy metabolism   |
| PA3661<br>PA3662 | hypothetical protein<br>hypothetical protein                                    | Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown                |
| PA3662<br>PA3674 | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA3682           | hypothetical protein  | Hypothetical, unclassified, unknown   |
|                  | adenylate kinase  | Nucleotide biosynthesis and metabolism  |
| PA3686           |   |   |
| PA3686<br>PA3692 | probable outer membrane protein precursor                                       | Membrane proteins   |
|                  | probable outer membrane protein precursor<br>probable transcriptional regulator | Membrane proteins<br>Transcriptional regulators   |

| ORF            | Product name   | Function class  |
|----------------|--|---|
| PA3718         | probable major facilitator superfamily (MFS) transporter | Membrane proteins   |
| PA3724         | elastase LasB  | Secreted Factors (toxins, enzymes, alginate)              |
| PA3737         | thiol:disulfide interchange protein DsbC                 | Translation, post-translational modification, degradation |
| PA3779         | hypothetical protein                                     | Hypothetical, unclassified, unknown                       |
| PA3780         | hypothetical protein                                     | Membrane proteins   |
| PA3781         | probable transporter                                     | Membrane proteins   |
| A3789          | hypothetical protein                                     | Membrane proteins   |
| PA3795         | probable oxidoreductase                                  | Putative enzymes  |
| PA3836         | hypothetical protein                                     | Hypothetical, unclassified, unknown                       |
| PA3837         | probable permease of ABC transporter                     | Membrane proteins   |
| PA3838         | probable ATP-binding component of ABC transporter        | Transport of small molecules                              |
| PA3839         | probable sodium:sulfate symporter                        | Membrane proteins   |
| PA3840         | conserved hypothetical protein                           | Hypothetical, unclassified, unknown                       |
| PA3865         | probable amino acid binding protein                      | Transport of small molecules                              |
| PA3880         | conserved hypothetical protein                           | Hypothetical, unclassified, unknown                       |
| A3887          | Na+/H+ antiporter NhaP                                   | Membrane proteins   |
| A3906          | hypothetical protein                                     | Hypothetical, unclassified, unknown                       |
|                |  |   |
| A3908          | hypothetical protein                                     | Hypothetical, unclassified, unknown                       |
| A3913          | probable protease  | Putative enzymes  |
| A3940          | probable DNA binding protein                             | DNA replication, recombination, modification and repair   |
| PA3959         | hypothetical protein                                     | Hypothetical, unclassified, unknown                       |
| PA3960         | hypothetical protein                                     | Hypothetical, unclassified, unknown                       |
| PA3966         | hypothetical protein                                     | Membrane proteins   |
| PA3967         | hypothetical protein                                     | Hypothetical, unclassified, unknown                       |
| PA3984         | apolipoprotein N-acyltransferase                         | Cell wall / LPS / capsule                                 |
| PA4021         | probable transcriptional regulator                       | Transcriptional regulators                                |
| PA4023         | probable transport protein                               | Transport of small molecules                              |
| PA4024         | ethanolamine ammonia-lyase large subunit                 | Central intermediary metabolism                           |
| PA4025         | probable ethanolamine ammonia-lyase light chain          | Carbon compound catabolism                                |
| PA4031         | inorganic pyrophosphatase                                | Central intermediary metabolism                           |
| PA4055         | riboflavin synthase alpha chain                          | Biosynthesis of cofactors, prosthetic groups and carriers |
| PA4057         | NrdR   | Nucleotide biosynthesis and metabolism                    |
| PA4058         | hypothetical protein                                     | Hypothetical, unclassified, unknown                       |
| PA4070         | probable transcriptional regulator                       | Transcriptional regulators                                |
| PA4072         | probable amino acid permease                             | Transport of small molecules                              |
| PA4073         | probable aldehyde dehydrogenase                          | Putative enzymes  |
| PA4078         | probable uneribosomal peptide synthetase                 | Adaptation, Protection                                    |
| PA4078         | probable dehydrogenase                                   | Putative enzymes  |
| PA4079         | 4-hydroxyphenylacetate 3-monooxygenase large chain       | Carbon compound catabolism                                |
|                | 4-hydroxyphenylacetate 3-monooxygenase small chain       | *   |
| PA4092         |  | Carbon compound catabolism                                |
| PA4093         | hypothetical protein                                     | Hypothetical, unclassified, unknown                       |
| PA4094         | probable transcriptional regulator                       | Transcriptional regulators                                |
| PA4120         | probable transcriptional regulator                       | Transcriptional regulators                                |
| PA4121         | conserved hypothetical protein                           | Hypothetical, unclassified, unknown                       |
| PA4122         | conserved hypothetical protein                           | Hypothetical, unclassified, unknown                       |
| PA4123         | 5-carboxy-2-hydroxymuconate semialdehyde dehydrogenase   | Carbon compound catabolism                                |
| PA4124         | homoprotocatechuate 2,3-dioxygenase                      | Carbon compound catabolism                                |
| PA4125         | 5-carboxymethyl-2-hydroxymuconate isomerase              | Carbon compound catabolism                                |
| PA4126         | probable major facilitator superfamily (MFS) transporter | Membrane proteins   |
| PA4127         | 2-oxo-hept-3-ene-1,7-dioate hydratase                    | Carbon compound catabolism                                |
| PA4128         | conserved hypothetical protein                           | Putative enzymes  |
| A4129          | hypothetical protein                                     | Hypothetical, unclassified, unknown                       |
| A4130          | probable sulfite or nitrite reductase                    | Central intermediary metabolism                           |
| A4131          | probable iron-sulfur protein                             | Putative enzymes  |
| A4132          | conserved hypothetical protein                           | Hypothetical, unclassified, unknown                       |
| A4133          | cytochrome c oxidase subunit (cbb3-type)                 | Energy metabolism   |
| A4134          | hypothetical protein                                     | Hypothetical, unclassified, unknown                       |
| A4136          | probable major facilitator superfamily (MFS) transporter | Membrane proteins   |
| A4137          | probable proin   | Transport of small molecules                              |
| A4137<br>A4140 | hypothetical protein                                     | Hypothetical, unclassified, unknown                       |
| A4140<br>A4141 | hypothetical protein                                     | Hypothetical, unclassified, unknown                       |
|                |  |   |
| PA4142         | probable secretion protein                               | Protein secretion/export apparatus                        |
| PA4147         | transcriptional regulator AcoR                           | Transcriptional regulators                                |
| PA4148         | probable short-chain dehydrogenase                       | Putative enzymes  |
| PA4150         | probable dehydrogenase E1 component                      | Carbon compound catabolism                                |
| PA4151         | acetoin catabolism protein AcoB                          | Carbon compound catabolism                                |
| PA4152         | probable hydrolase                                       | Carbon compound catabolism                                |
| PA4153         | 2,3-butanediol dehydrogenase                             | Carbon compound catabolism                                |

| ORF              | Product name  | Function class  |
|------------------|---|---|
| PA4175           | protease IV   | Secreted Factors (toxins, enzymes, alginate)                                  |
| PA4180           | probable acetolactate synthase large subunit  | Amino acid biosynthesis and metabolism  |
| PA4205           | hypothetical protein  | Membrane proteins   |
| PA4206           | probable Resistance-Nodulation-Cell Division (RND) efflux membrane fusion protein         |   |
| PA4207           | probable Resistance-Nodulation-Cell Division (RND) efflux transporter                     | Membrane proteins   |
| PA4208           | probable outer membrane protein precursor   | Membrane proteins   |
| PA4209           | probable phenazine-specific methyltransferase   | Putative enzymes  |
| PA4210           | probable phenazine biosynthesis protein   | Secreted Factors (toxins, enzymes, alginate)                                  |
| PA4211           | probable phenazine biosynthesis protein   | Secreted Factors (toxins, enzymes, alginate)                                  |
| PA4288           | probable transcriptional regulator  | Transcriptional regulators  |
| PA4293           | two-component sensor PprA   | Two-component regulatory systems  |
| PA4294           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA4297           | TadG  | Membrane proteins   |
| PA4298           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA4316           | exodeoxyribonuclease I  | DNA replication, recombination, modification and repair                       |
| PA4324           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA4328           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA4329           | pyruvate kinase II  | Energy metabolism   |
| PA4348           | conserved hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA4351           | probable acyltransferase  | Fatty acid and phospholipid metabolism  |
| PA4352           | conserved hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA4357           | conserved hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA4358<br>PA4377 | probable ferrous iron transport protein<br>hypothetical protein                           | Membrane proteins<br>Hypothetical, unclassified, unknown                      |
| PA4394           | conserved hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA4433           | 50S ribosomal protein L13   | Translation, post-translational modification, degradation                     |
| PA4436           | probable transcriptional regulator  | Transcriptional regulators  |
| PA4496           | probable binding protein component of ABC transporter                                     | Transport of small molecules  |
| PA4497           | probable binding protein component of ABC transporter                                     | Transport of small molecules  |
| PA4498           | probable metallopeptidase   | Translation, post-translational modification, degradation                     |
| PA4500           | probable binding protein component of ABC transporter                                     | Transport of small molecules  |
| PA4502           | probable binding protein component of ABC transporter                                     | Transport of small molecules  |
| PA4503           | probable permease of ABC transporter  | Membrane proteins   |
| PA4504           | probable permease of ABC transporter  | Membrane proteins   |
| PA4505           | probable ATP-binding component of ABC transporter   | Transport of small molecules  |
| PA4506           | probable ATP-binding component of ABC dipeptide transporter                               | Transport of small molecules  |
| PA4520           | probable chemotaxis transducer  | Adaptation, Protection  |
| PA4569<br>PA4577 | octaprenyl-diphosphate synthase<br>hypothetical protein                                   | Energy metabolism<br>Hypothetical, unclassified, unknown                      |
| PA4588           | glutamate dehydrogenase   | Amino acid biosynthesis and metabolism  |
| PA4596           | probable transcriptional regulator  | Transcriptional regulators  |
| PA4606           | conserved hypothetical protein  | Adaptation, Protection  |
| PA4611           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA4632           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA4633           | probable chemotaxis transducer  | Adaptation, Protection  |
| PA4654           | probable major facilitator superfamily (MFS) transporter                                  | Membrane proteins   |
| PA4676           | probable carbonic anhydrase   | Putative enzymes  |
| PA4677           |   | Hypothetical, unclassified, unknown   |
| PA4681           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA4682           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA4683           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA4695<br>PA4715 | acetolactate synthase isozyme III small subunit<br>probable aminotransferase              | Biosynthesis of cofactors, prosthetic groups and carriers<br>Putative enzymes |
| PA4713<br>PA4733 | acetyl-coenzyme A synthetase  | Carbon compound catabolism  |
| PA4757           | conserved hypothetical protein  | Membrane proteins   |
| PA4758           | carbamoyl-phosphate synthase small chain  | Amino acid biosynthesis and metabolism  |
| PA4759           | dihydrodipicolinate reductase   | Amino acid biosynthesis and metabolism  |
| PA4770           | L-lactate permease  | Transport of small molecules  |
| PA4773           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA4779           | hypothetical protein  | Membrane proteins   |
| PA4782           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA4808           | L-seryl-tRNA(ser) selenium transferase  | Translation, post-translational modification, degradation                     |
| PA4817           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA4851           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA4852           | conserved hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA4860<br>PA4862 | probable permease of ABC transporter<br>probable ATP-binding component of ABC transporter | Membrane proteins<br>Transport of small molecules                             |
| PA4862<br>PA4867 | urease beta subunit   | Central intermediary metabolism   |
| 111100/          |   | contai mermenary memorinim  |

| ORF              | Product name  | Function class  |
|------------------|---|---|
| PA4884           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA4893           | urease accessory protein UreG   | Biosynthesis of cofactors, prosthetic groups and carriers                             |
| PA4894           | hypothetical protein  | Membrane proteins   |
| PA4899           | probable aldehyde dehydrogenase   | Putative enzymes  |
| PA4901           | benzoylformate decarboxylase  | Carbon compound catabolism  |
| PA4908<br>PA4910 | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA4910<br>PA4911 | probable ATP-binding component of ABC transporter<br>probable permease of ABC branched-chain amino acid transporter | Transport of small molecules<br>Membrane proteins                                     |
| PA4912           | probable permease of ABC branched chain amino acid transporter  | Membrane proteins   |
| PA4913           | probable binding protein component of ABC transporter   | Transport of small molecules  |
| PA4943           | probable GTP-binding protein  | Putative enzymes  |
| PA4979           | probable acyl-CoA dehydrogenase   | Putative enzymes  |
| PA4983           | probable two-component response regulator   | Transcriptional regulators  |
| PA4984           | probable transcriptional regulator  | Transcriptional regulators  |
| PA4985           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA4986           | probable oxidoreductase   | Putative enzymes  |
| PA4997<br>PA5020 | transport protein MsbA<br>probable acyl-CoA dehydrogenase   | Fatty acid and phospholipid metabolism<br>Putative enzymes                            |
| PA5020<br>PA5025 | homocysteine synthase   | Amino acid biosynthesis and metabolism  |
| PA5025<br>PA5027 | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA5028           | conserved hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA5030           | probable major facilitator superfamily (MFS) transporter  | Membrane proteins   |
| PA5051           | arginyl-tRNA synthetase   | Translation, post-translational modification, degradation                             |
| PA5059           | probable transcriptional regulator  | Transcriptional regulators  |
| PA5082           | probable binding protein component of ABC transporter   | Transport of small molecules  |
| PA5096           | probable binding protein component of ABC transporter   | Transport of small molecules  |
| PA5097           | probable amino acid permease  | Membrane proteins   |
| PA5098           | histidine ammonia-lyase   | Amino acid biosynthesis and metabolism  |
| PA5106<br>PA5112 | conserved hypothetical protein<br>esterase EstA   | Hypothetical, unclassified, unknown<br>Secreted Factors (toxins, enzymes, alginate)   |
| PA5129           | glutaredoxin  | Energy metabolism   |
| PA5130           | conserved hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA5137           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA5149           | conserved hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA5152           | probable ATP-binding component of ABC transporter   | Transport of small molecules  |
| PA5153           | probable periplasmic binding protein  | Transport of small molecules  |
| PA5154           | probable permease of ABC transporter  | Membrane proteins   |
| PA5155           | probable permease of ABC transporter  | Membrane proteins   |
| PA5161<br>PA5162 | dTDP-D-glucose 4,6-dehydratase<br>dTDP-4-dehydrorhamnose reductase  | Carbon compound catabolism  |
| PA5162<br>PA5163 | glucose-1-phosphate thymidylyltransferase   | Cell wall / LPS / capsule<br>Cell wall / LPS / capsule                                |
| PA5164           | dTDP-4-dehydrorhamnose 3,5-epimerase  | Cell wall / LPS / capsule   |
| PA5168           | probable dicarboxylate transporter  | Membrane proteins   |
| PA5169           | probable C4-dicarboxylate transporter   | Membrane proteins   |
| PA5174           | probable beta-ketoacyl synthase   | Fatty acid and phospholipid metabolism  |
| PA5180           | conserved hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA5220           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA5237           | conserved hypothetical protein<br>thioredoxin   | Hypothetical, unclassified, unknown   |
| PA5240<br>PA5249 | hypothetical protein  | Translation, post-translational modification, degradation<br>Membrane proteins        |
| PA5249<br>PA5275 | conserved hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA5285           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA5286           | conserved hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA5288           | nitrogen regulatory protein P-II 2  | Central intermediary metabolism   |
| PA5289           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA5297           | pyruvate dehydrogenase (cytochrome)   | Central intermediary metabolism   |
| PA5303           | conserved hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA5309           | probable oxidoreductase   | Putative enzymes  |
| PA5324           | probable transcriptional regulator  | Transcriptional regulators  |
| PA5325<br>PA5328 | hypothetical protein<br>probable cytochrome c(mono-heme type)   | Hypothetical, unclassified, unknown<br>Energy metabolism                              |
| PA5328<br>PA5329 | conserved hypothetical protein  | Energy metabolism<br>Hypothetical, unclassified, unknown                              |
| PA5329<br>PA5338 | guanosine-3',5'-bis(diphosphate) 3'-pyrophosphohydrolase  | Nucleotide biosynthesis and metabolism  |
| PA5344           | probable transcriptional regulator  | Transcriptional regulators  |
|                  |   |   |
| PA5348           | probable DNA-binding protein  | DNA replication, recombination, modification and repair                               |
|                  | · · ·   | DNA replication, recombination, modification and repair<br>Carbon compound catabolism |
| PA5348           | probable DNA-binding protein  |   |

| ORF    | Product name  | Function class                      |
|--------|---|-------------------------------------|
|        |   |                                     |
| PA5380 | GbdR  | Transcriptional regulators          |
| PA5381 | hypothetical protein                                      | Hypothetical, unclassified, unknown |
| PA5400 | probable electron transfer flavoprotein alpha subunit     | Energy metabolism                   |
| PA5409 | hypothetical protein                                      | Membrane proteins                   |
| PA5427 | alcohol dehydrogenase                                     | Energy metabolism                   |
| PA5434 | tryptophan permease                                       | Membrane proteins                   |
| PA5435 | probable transcarboxylase subunit                         | Central intermediary metabolism     |
| PA5436 | probable biotin carboxylase subunit of a transcarboxylase | Central intermediary metabolism     |
| PA5442 | conserved hypothetical protein                            | Membrane proteins                   |
| PA5445 | probable coenzyme A transferase                           | Putative enzymes                    |
| PA5451 | membrane subunit of A-band LPS efflux transporter         | Cell wall / LPS / capsule           |
| PA5452 | phosphomannose isomerase/GDP-mannose WbpW                 | Cell wall / LPS / capsule           |
| PA5461 | hypothetical protein                                      | Hypothetical, unclassified, unknown |
| PA5475 | hypothetical protein                                      | Hypothetical, unclassified, unknown |
| PA5491 | probable cytochrome                                       | Energy metabolism                   |
| PA5506 | hypothetical protein                                      | Hypothetical, unclassified, unknown |
| PA5507 | hypothetical protein                                      | Hypothetical, unclassified, unknown |
| PA5508 | probable glutamine synthetase                             | Putative enzymes                    |
| PA5509 | hypothetical protein                                      | Hypothetical, unclassified, unknown |
| PA5510 | probable transporter                                      | Membrane proteins                   |
| PA5522 | probable glutamine synthetase                             | Putative enzymes                    |
| PA5523 | probable aminotransferase                                 | Putative enzymes                    |
| PA5533 | hypothetical protein                                      | Hypothetical, unclassified, unknown |
| PA5536 | conserved hypothetical protein                            | Hypothetical, unclassified, unknown |
| PA5542 | hypothetical protein                                      | Putative enzymes                    |
| PA5543 | hypothetical protein                                      | Hypothetical, unclassified, unknown |
| PA5544 | conserved hypothetical protein                            | Membrane proteins                   |
| PA5545 | conserved hypothetical protein                            | Hypothetical, unclassified, unknown |
| PA5546 | conserved hypothetical protein                            | Putative enzymes                    |

### ORF Product name

#### **Oxidative stress**

| PA0007hypothetical proteinPA0005conserved hypothetical proteinPA0105cytochrome c oxidase, subunit IPA0107conserved hypothetical proteinPA0108conserved hypothetical proteinPA0128conserved hypothetical proteinPA0128conserved hypothetical proteinPA0171hypothetical proteinPA0173probable methylesterasePA0174conserved hypothetical proteinPA0175aerotaxis transducer Aer2PA0176aerotaxis transducer Aer2PA0177probable two-component sensorPA0178probable two-component response regulatorPA0179probable two-component response regulatorPA0280sulfate transport protein CysXPA0281sulfate transport protein CysWPA0282sulfate transport protein CysTPA0283sulfate transport protein CysTPA0335phypothetical proteinPA0352probable biotin-requiring enzymePA04353phypothetical proteinPA0454probable ClpA/B protein MicAPA0455fructose-1, 6-bisphosphate aldolasePA0556fructose-1, 6-bisphosphate aldolasePA0557fructose-1, 6-bisphosphate aldolasePA0588ohypothetical proteinPA0556probable taeriophage proteinPA0557ructose-1, 6-bisphosphate aldolasePA0558conserved hypothetical proteinPA0559phypothetical proteinPA0550probable bacteriophage proteinPA0517probable bacteriophage protein </th <th></th> <th></th>                             |        |   |
|--|--------|---|
| PA0105cytochrome c oxidase, subunit IIPA0107conserved hypothetical proteinPA0107conserved hypothetical proteinPA0128conserved hypothetical proteinPA0128conserved hypothetical proteinPA0171hypothetical proteinPA0173probable methylesterasePA0174conserved hypothetical proteinPA0175aerotaxis transducer Aer2PA0176aerotaxis transducer Aer2PA0177probable two-component sensorPA0178probable two-component sensorPA0182probable two-component sensorPA0183sulfate transport protein CysAPA0280sulfate transport protein CysAPA0281sulfate transport protein CysTPA0382sulfate transport protein protein CysTPA0383hypothetical proteinPA0384probable biotin-requiring enzymePA0459probable lotin-requiring enzymePA0459probable lotin-requiring enzymePA0459probable lotin-requiring enzymePA0459probable lotin-requiring enzymePA0455fructose-1, 6-bisphosphate aldolasePA0555fructose-1, 6-bisphosphate aldolasePA0561repressor, PtrBPA0617probable bacteriophage proteinPA0618probable bacteriophage proteinPA0619probable bacteriophage proteinPA0616hypothetical proteinPA0617probable bacteriophage proteinPA0618probable bacteriophage proteinPA0619probable bacteriophage protein <td></td> <td>hypothetical protein</td>                    |        | hypothetical protein                                |
| PA0106cytochrome c oxidase, subunit IPA0109conserved hypothetical proteinPA0122conserved hypothetical proteinPA0123conserved hypothetical proteinPA0124conserved hypothetical proteinPA0175probable methylesterasePA0171hypothetical proteinPA0173probable methylesterasePA0174conserved hypothetical proteinPA0175acrotaxis transducer Aer2PA0176acrotaxis transducer Aer2PA0177probable two-component response regulatorPA0125probable two-component response regulatorPA0126probable two-component response regulatorPA0127probable two-component response regulatorPA0128sulfate transport protein CysAPA0280sulfate transport protein CysAPA0281sulfate transport protein CysAPA0282sulfate transport protein CysAPA0283sulfate transport protein CysAPA0385hypothetical proteinPA0385hypothetical proteinPA0385hypothetical proteinPA0493probable typothetical proteinPA0493probable solution requiring enzymePA0494probable solution requiring enzymePA0495fructose-1,6-bisphosphate aldolasePA0555fructose-1,6-bisphosphate aldolasePA0555fructose-1,6-bisphosphate aldolasePA0555fructose-1,6-bisphosphate aldolasePA0555rpoteinPA0612repressor, PtrBPA0613hypothetical protein <tr< td=""><td></td><td></td></tr<>                             |        |   |
| PA0107conserved hypothetical proteinPA0128conserved hypothetical proteinPA0128conserved hypothetical proteinPA0158RND; triclosan efflux transporter, TriCPA01711hypothetical proteinPA0173probable methylesterasePA0174conserved hypothetical proteinPA0175aerotaxis transducer Aer2PA0177probable two-component sensorPA0178probable two-component response regulatorPA0179probable two-component response regulatorPA0182probable short-chain dehydrogenasePA0280sulfate transport protein CysAPA0281sulfate transport protein CysTPA0283sulfate transport protein CysTPA0284sulfate transport protein CysTPA0355phospheonolpyruvate-protein phosphotransferase PtsPPA0352probable transportPA0385hypothetical proteinPA0439probable tonin-requiring enzymePA0439probable botin-requiring enzymePA0439probable acyl-CoA carboxylase subunitPA0492conserved hypothetical proteinPA0493probable bacteriophage proteinPA0494conserved hypothetical proteinPA0556fructose-1, 6-bisphosphate aldolasePA0586conserved hypothetical proteinPA0587conserved hypothetical proteinPA0618hypothetical proteinPA0619probable bacteriophage proteinPA0614hypothetical proteinPA0615hypothetical proteinPA0626pro  |        |   |
| PA0109hypothetical proteinPA0122conserved hypothetical proteinPA0128conserved hypothetical proteinPA0171hypothetical proteinPA0173probable methylesterasePA0174conserved hypothetical proteinPA0175aerotaxis transducer Aer2PA0177probable prine-binding chemotaxis proteinPA0178aerotaxis transducer Aer2PA0179probable two-component sensorPA0182probable two-component sensorPA0183probable two-component sensorPA0184probable short-chain dehydrogenasePA0280sulfate transport protein CysAPA0281sulfate transport protein CysTPA0283sulfate transport protein precursorPA0383hybothetical proteinPA0383hybothetical proteinPA0384probable transporterPA0385hypothetical proteinPA0495probable doil-CoA carboxylase subunitPA0496probable doil-CoA carboxylase subunitPA0497probable doil-coveliaPA0498probable doil-coveliaPA0499probable doil-coveliaPA0491probable doil-coveliaPA0492conserved hypothetical proteinPA0493probable aproteinPA0586conserved hypothetical proteinPA0587conserved hypothetical proteinPA0586conserved hypothetical proteinPA0617probable bacteriophage proteinPA0618probable bacteriophage proteinPA0619probable bacteriophage protein<   |        |   |
| PA0122conserved hypothetical proteinPA0158conserved hypothetical proteinPA0158RND; triclosan efflux transporter, TriCPA0171hypothetical proteinPA0173probable methylesterasePA0174conserved hypothetical proteinPA0175aerotaxis transducer Aer2PA0176aerotaxis transducer Aer2PA0177probable two-component response regulatorPA0182probable two-component response regulatorPA0193probable two-component response regulatorPA0194probable short-chain dehydrogenasePA02950conserved hypothetical proteinPA0281sulfate transport protein CysAPA0282sulfate transport protein CysTPA0283sulfate-binding protein precursorPA0337phosphoenolpyruvate-protein MicAPA0337phosphoenolpyruvate-protein MicAPA0382DNA mismatch repair protein MicAPA0439probable ClpA/B protease ATP binding subunitPA0430probable ClpA/B protease ATP binding subunitPA0431probable clpA/B protease ATP binding subunitPA0432conserved hypothetical proteinPA0433probable acyl-CoA carboxylase subunitPA0434probable acyl-CoA carboxylase subunitPA0435hypothetical proteinPA0556fructose-1, 6-bisphosphate aldolasePA0557fructose-1, 6-bisphosphate aldolasePA0586conserved hypothetical proteinPA0611hypothetical proteinPA0612nproteinPA0613hypotheti                         |        |   |
| PA0128conserved hypothetical proteinPA0138RND; triclosan efflux transporter, TriCPA0171hypothetical proteinPA0173probable methylesterasePA0174conserved hypothetical proteinPA0175probable purine-binding chemotaxis proteinPA0176aerotaxis transducer Aer2PA0177probable two-component response regulatorPA0178probable short-chain dehydrogenasePA0129probable short-chain dehydrogenasePA0200conserved hypothetical proteinPA0230sulfate transport protein CysAPA0281sulfate transport protein CysTPA0335hypothetical proteinPA0335hypothetical protein precursorPA0335probable transport protein of MicAPA0382DNA mismatch repair protein MicAPA0433probable ClpA/B protease ATP binding subunitPA0434probable ColpA/B protease ATP binding subunitPA0435fructose-1,6-bisphosphate aldolasePA0555fructose-1,6-bisphosphate aldolasePA0582dihydroneopterin aldolasePA0583conserved hypothetical proteinPA0613hypothetical proteinPA0614hypothetical proteinPA0615hypothetical proteinPA0616hypothetical proteinPA0617probable bacteriophage proteinPA0618probable bacteriophage proteinPA0619probable bacteriophage proteinPA0620probable bacteriophage proteinPA0614hypothetical proteinPA0625pro  |        |   |
| PA0158RND; triclosan efflux transporter, TriCPA0171hypothetical proteinPA0173probable methylesterasePA0174conserved hypothetical proteinPA0176aerotaxis transducer Aer2PA0177probable two-component sensorPA0178probable two-component sensorPA0179probable two-component sensorPA0182probable short-chain dehydrogenasePA0200conserved hypothetical proteinPA0281sulfate transport protein CysAPA0282sulfate transport protein CysTPA0335hypothetical proteinPA0337phosphoenolpyruvate-protein phosphotransferase PtsPPA0338probable ClpA/B protease ATP binding subunitPA0439probable ClpA/B protease ATP binding subunitPA0439probable ClpA/B protease ATP binding subunitPA0439probable biotin-requiring enzymePA0431probable acyl-CoA carboxylase subunitPA0432conserved hypothetical proteinPA055fructose-1,6-bisphosphate aldolasePA052conserved hypothetical proteinPA0511hypothetical proteinPA0522conserved hypothetical proteinPA0531hypothetical proteinPA0542dihydroneopterin aldolasePA0552probable bacteriophage proteinPA0563hypothetical proteinPA0512repressor, PtrBPA0613hypothetical proteinPA0614hypothetical proteinPA0615hypothetical proteinPA0622probable bacteriophage prot  |        | 51 1  |
| PA0171hypothetical proteinPA0173probable methylesterasePA0174conserved hypothetical proteinPA0176aerotaxis transducer Aer2PA0177probable buvo-component response regulatorPA0178probable two-component response regulatorPA0182probable short-chain dehydrogenasePA0280sulfate transport protein CysAPA0281sulfate transport protein CysAPA0282sulfate transport protein CysTPA0283sulfate-binding protein precursorPA0335hypothetical proteinPA0337phosphoenolpyruvate-protein hhosphotransferase PtsPPA0332probable ClpA/B protease ATP binding subunitPA0492conserved hypothetical proteinPA0493probable ClpA/B protease ATP binding subunitPA0494probable ClpA/B protease ATP binding subunitPA0495hypothetical proteinPA0496conserved hypothetical proteinPA0497probable acyl-CoA carboxylase subunitPA0498probable acyl-CoA carboxylase subunitPA0499probable acyl-CoA carboxylase subunitPA0491probable acyl-CoA carboxylase subunitPA0492conserved hypothetical proteinPA0586conserved hypothetical proteinPA0587conserved hypothetical proteinPA0588dihydroneopterin aldolasePA0512repressor, PtrBPA0613hypothetical proteinPA0614hypothetical proteinPA0615hypothetical proteinPA0616hypothetical protein <tr< td=""><td></td><td></td></tr<>  |        |   |
| PA0173probable methylesterasePA0174conserved hypothetical proteinPA0176aerotaxis transducer Aer2PA0177probable purine-binding chemotaxis proteinPA0178probable two-component response regulatorPA0179probable short-chain dehydrogenasePA0120conserved hypothetical proteinPA0220sulfate transport protein CysAPA0281sulfate transport protein CysTPA0283sulfate transport protein presensorPA0335hypothetical proteinPA0337phosphoenolpyruvate-protein phosphotransferase PtsPPA0338probable transporterPA0385hypothetical proteinPA0494probable ClpA/B protease ATP binding subunitPA0495probable ClpA/B protease ATP binding subunitPA0496probable acyl-CoA carboxylase subunitPA0497hypothetical proteinPA0498probable acyl-CoA carboxylase subunitPA0495hypothetical proteinPA0556conserved hypothetical proteinPA0613hypothetical proteinPA0614hypothetical proteinPA0615hypothetical proteinPA0616hypothetical proteinPA0617probable bacteriophage proteinPA0618probable bacteriophage proteinPA0619probable bacteriophage proteinPA0614hypothetical proteinPA0625hypothetical proteinPA0626probable bacteriophage proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical pro  |        |   |
| PA0176aerotaxis transducer Aer2PA0177probable purine-binding chemotaxis proteinPA0178probable two-component response regulatorPA0179probable two-component response regulatorPA0182probable short-chain dehydrogenasePA0280sulfate transport protein CysAPA0281sulfate transport protein CysYPA0282sulfate transport protein CysYPA0283sulfate-binding protein precursorPA0335hypothetical proteinPA0335phosphoenolpyruvate-protein hhosphotransferase PtsPPA0335probable ClpA/B protease ATP binding subunitPA0459probable ClpA/B protease ATP binding subunitPA0459probable cayl-CoA carboxylase subunitPA0459probable acyl-CoA carboxylase subunitPA0459probable acyl-CoA carboxylase subunitPA0451hypothetical proteinPA0555fructose-1,6-bisphosphate aldolasePA0586conserved hypothetical proteinPA0612repressor, PtrBPA0613hypothetical proteinPA0614hypothetical proteinPA0615probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0644hypothetical proteinPA0655hypothetical proteinPA0626hypothetical proteinPA0627probable bacteriophage proteinPA0628conserved hypothetical protein </td <td>PA0173</td> <td></td> | PA0173 |   |
| PA0177probable purine-binding chemotaxis proteinPA0178probable two-component response regulatorPA0179probable short-chain dehydrogenasePA0250conserved hypothetical proteinPA0281sulfate transport protein CysAPA0282sulfate transport protein CysTPA0283sulfate transport protein precursorPA0335hypothetical proteinPA0337phosphoenolpyruvate-protein micAPA0338probable transporterPA0339probable ClpA/B protease ATP binding subunitPA0459probable ClpA/B protease ATP binding subunitPA0459probable clpA/B protease ATP binding subunitPA0492conserved hypothetical proteinPA0493probable acyl-CoA carboxylase subunitPA0494probable acyl-CoA carboxylase subunitPA0455fructose-1,6-bisphosphate alolasePA0586conserved hypothetical proteinPA0587conserved hypothetical proteinPA0618hypothetical proteinPA0519probable bacteriophage proteinPA0612repressor, PtrBPA0613hypothetical proteinPA0614hypothetical proteinPA0615hypothetical proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0619probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622hypothetical proteinPA0623   | PA0174 | conserved hypothetical protein                      |
| PA0178probable two-component sensorPA0179probable two-component response regulatorPA0182probable short-chain dehydrogenasePA0250conserved hypothetical proteinPA0281sulfate transport protein CysKPA0282sulfate transport protein CysTPA0283sulfate transport protein CysTPA0284sulfate transport protein precursorPA0335hypothetical proteinPA0337phosphoenolpyruvate-protein MicAPA0382DNA mismatch repair protein MicAPA0383hypothetical proteinPA0384probable ClpA/B protease ATP binding subunitPA0459probable ClpA/B protease ATP binding subunitPA0492conserved hypothetical proteinPA0493probable acyl-CoA carboxylase subunitPA0494probable acyl-CoA carboxylase subunitPA0585fructose-1,6-bisphosphate aldolasePA0586conserved hypothetical proteinPA0587conserved hypothetical proteinPA0613hypothetical proteinPA0614hypothetical proteinPA0615hypothetical proteinPA0616hypothetical proteinPA0617probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetic  | PA0176 | aerotaxis transducer Aer2                           |
| PA0179probable two-component response regulatorPA0182probable short-chain dehydrogenasePA0280conserved hypothetical proteinPA0281sulfate transport protein CysMPA0282sulfate transport protein CysTPA0283sulfate transport protein CysTPA0284sulfate transport protein CysTPA0335hypothetical protein precursorPA0335hypothetical protein phosphotransferase PtsPPA0332probable transporterPA0333hypothetical proteinPA0459probable ClpA/B protease ATP binding subunitPA0459probable ClpA/B protease ATP binding subunitPA0492conserved hypothetical proteinPA0493probable col-CoA carboxylase subunitPA0494probable acyl-CoA carboxylase subunitPA0495hypothetical proteinPA0555fructose-1, 6-bisphosphate aldolasePA0586conserved hypothetical proteinPA0587conserved hypothetical proteinPA0613hypothetical proteinPA0614hypothetical proteinPA0615hypothetical proteinPA0616hypothetical proteinPA0617probable bacteriophage proteinPA0628conserved hypothetical proteinPA0619probable bacteriophage proteinPA0620probable bacteriophage proteinPA0616hypothetical proteinPA0620probable bacteriophage proteinPA0621probable bacteriophage proteinPA0622probable bacteriophage proteinPA0623prob  |        |   |
| PA0182probable short-chain dehydrogenasePA0250conserved hypothetical proteinPA0280sulfate transport protein CysWPA0281sulfate transport protein CysWPA0282sulfate transport protein precursorPA0333hypothetical proteinPA0337phosphoenolpyruvate-protein phosphotransferase PtsPPA0338probable transporterPA0382DNA mismatch repair protein MicAPA0459probable ClpA/B protease ATP binding subunitPA0459probable ClpA/B protease ATP binding subunitPA0493probable clpA/B protease ATP binding subunitPA0494probable acyl-CoA carboxylase subunitPA0495hypothetical proteinPA0496hypothetical proteinPA0555fructose-1,6-bisphosphate aldolasePA0586conserved hypothetical proteinPA0611hypothetical proteinPA0612repressor, PtrBPA0613hypothetical proteinPA0614hypothetical proteinPA0615hypothetical proteinPA0616hypothetical proteinPA0617probable bacteriophage proteinPA0628conserved hypothetical proteinPA0618probable bacteriophage proteinPA0629probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical protein <td></td> <td></td>   |        |   |
| PA0250conserved hypothetical proteinPA0280sulfate transport protein CysAPA0281sulfate transport protein CysTPA0282sulfate transport protein precursorPA0335hypothetical protein precursorPA0337phosphoenolpyruvate-protein phosphotransferase PtsPPA0352probable transporterPA0385hypothetical proteinPA0385hypothetical proteinPA0459probable ClpA/B protease ATP binding subunitPA0459probable ClpA/B protease ATP binding subunitPA0492conserved hypothetical proteinPA0493probable acyl-CoA carboxylase subunitPA0494probable acyl-CoA carboxylase subunitPA0495hypothetical proteinPA0555fructose-1,6-bisphosphate aldolasePA0582dihydroneopterin aldolasePA0586conserved hypothetical proteinPA0611hypothetical proteinPA0612repressor, PtrBPA0613hypothetical proteinPA0614hypothetical proteinPA0615hypothetical proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0639probable bacteriophage proteinPA0620 <td></td> <td></td>   |        |   |
| PA0280sulfate transport protein CysAPA0281sulfate transport protein CysWPA0282sulfate transport protein CysTPA0283sulfate-binding protein precursorPA0335hypothetical proteinPA0337phosphoenolpyruvate-protein phosphotransferase PtsPPA0332probable transporterPA0382DNA mismatch repair protein MicAPA0459probable ClpA/B protease ATP binding subunitPA0459probable ClpA/B protease ATP binding subunitPA0492conserved hypothetical proteinPA0493probable col-CoA carboxylase subunitPA0494probable acyl-CoA carboxylase subunitPA0495hypothetical proteinPA0556fructose-16-bisphosphate aldolasePA0586conserved hypothetical proteinPA0612repressor, PtrBPA0613hypothetical proteinPA0614hypothetical proteinPA0615hypothetical proteinPA0616hypothetical proteinPA0617probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0639hypothetical proteinPA0630hypothetical proteinPA0631h   |        |   |
| PA0281sulfate transport protein CysWPA0282sulfate transport protein CysTPA0283sulfate-binding protein precursorPA0335hypothetical proteinPA0335phosphoenolpytuvate-protein phosphotransferase PtsPPA0352probable transporterPA0385hypothetical proteinPA0439probable ClpA/B protease ATP binding subunitPA0490probable ClpA/B protease ATP binding subunitPA0491probable ClpA/B protease ATP binding subunitPA0492conserved hypothetical proteinPA0493probable dayl-CoA carboxylase subunitPA0494probable acyl-CoA carboxylase subunitPA0495hypothetical proteinPA0585fructose-1,6-bisphosphate aldolasePA0586conserved hypothetical proteinPA0587conserved hypothetical proteinPA0612repressor, PtrBPA0613hypothetical proteinPA0616hypothetical proteinPA0617probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0639hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA  |        |   |
| PA0282sulfate transport protein CysTPA0283sulfate-binding protein precursorPA0335hypothetical proteinPA0337phosphoenolpyruvate-protein phosphotransferase PtsPPA0382DNA mismatch repair protein MicAPA0383hypothetical proteinPA0459probable LTpA/B protease ATP binding subunitPA0459probable ClpA/B protease ATP binding subunitPA0492conserved hypothetical proteinPA0493probable acyl-CoA carboxylase subunitPA0494probable acyl-CoA carboxylase subunitPA0495hypothetical proteinPA0555fructose-1,6-bisphosphate aldolasePA0582dihydroneopterin aldolasePA0586conserved hypothetical proteinPA0612repressor, PtrBPA0613hypothetical proteinPA0614hypothetical proteinPA0615hypothetical proteinPA0616hypothetical proteinPA0617probable bacteriophage proteinPA0618probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0639hypothetical proteinPA0630hypothetical proteinPA0631hypo   |        |   |
| PA0283sulfate-binding protein precursorPA0335hypothetical proteinPA0337phosphoenolpyruvate-protein phosphotransferase PtsPPA0332probable transporterPA0385hypothetical proteinPA0385hypothetical proteinPA0459probable ClpA/B protease ATP binding subunitPA0459probable binin-requiring enzymePA0493probable binin-requiring enzymePA0494probable acyl-CoA carboxylase subunitPA0495hypothetical proteinPA0496hypothetical proteinPA0555fructose-1,6-bisphosphate aldolasePA0586conserved hypothetical proteinPA0587conserved hypothetical proteinPA0613hypothetical proteinPA0614hypothetical proteinPA0615hypothetical proteinPA0616hypothetical proteinPA0617probable bacteriophage proteinPA0618probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0638probable bacteriophage proteinPA0639hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0626hypothetical proteinPA0627<   |        |   |
| PA0335hypothetical proteinPA0337phosphoenolpyruvate-protein phosphotransferase PtsPPA0352probable transporterPA0382DNA mismatch repair protein MicAPA0383hypothetical proteinPA0439probable ClpA/B protease ATP binding subunitPA0492conserved hypothetical proteinPA0493probable ClpA/B protease ATP binding subunitPA0494probable acyl-CoA carboxylase subunitPA0495hypothetical proteinPA0555fructose-1,6-bisphosphate aldolasePA0586conserved hypothetical proteinPA0587conserved hypothetical proteinPA0612repressor, PtrBPA0613hypothetical proteinPA0614hypothetical proteinPA0615hypothetical proteinPA0616hypothetical proteinPA0617probable bacteriophage proteinPA0618probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621proserved hypothetical proteinPA0622probable bacteriophage proteinPA0633probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0629conserved hypothetical proteinPA0620probable bacteriophage proteinPA0621probable bacteriophage proteinPA0622probable bacteriophagePA0633hypothe   |        |   |
| PA0352probable transporterPA0382DNA mismatch repair protein MicAPA0385hypothetical proteinPA0439probable CIpA/B protease ATP binding subunitPA0491conserved hypothetical proteinPA0492conserved hypothetical proteinPA0493probable biotin-requiring enzymePA0494probable acyl-CoA carboxylase subunitPA0495hypothetical proteinPA0555fructose-1,6-bisphosphate aldolasePA0582dihydroneopterin aldolasePA0583conserved hypothetical proteinPA0614hypothetical proteinPA0615hypothetical proteinPA0616hypothetical proteinPA0617probable bacteriophage proteinPA0618probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0639probable bacteriophage proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0638conserved hypothetical proteinPA0631hypothetical proteinPA06   | PA0335 |   |
| PA0382DNA mismatch repair protein MicAPA0385hypothetical proteinPA0459probable ClpA/B protease ATP binding subunitPA0492conserved hypothetical proteinPA0493probable biotin-requiring enzymePA0494probable acyl-CoA carboxylase subunitPA0495hypothetical proteinPA0555fructose-1,6-bisphosphate aldolasePA0582dihydroneopterin aldolasePA0583conserved hypothetical proteinPA0584conserved hypothetical proteinPA0612repressor, PtrBPA0613hypothetical proteinPA0614hypothetical proteinPA0615hypothetical proteinPA0616hypothetical proteinPA0617probable bacteriophage proteinPA0618probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0638conserved hypothetical proteinPA0639hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0632probable bacteriophage proteinPA0633probable bacteriophage proteinPA0641hypothetical proteinPA0630hypothetical proteinPA0631hypo  | PA0337 | phosphoenolpyruvate-protein phosphotransferase PtsP |
| PA0385hypothetical proteinPA0459probable ClpA/B protease ATP binding subunitPA0492conserved hypothetical proteinPA0493probable biotin-requiring enzymePA0494probable acyl-CoA carboxylase subunitPA0495hypothetical proteinPA0555fructose-1,6-bisphosphate aldolasePA0582dihydroneopterin aldolasePA0583conserved hypothetical proteinPA0612repressor, PtrBPA0613hypothetical proteinPA0614hypothetical proteinPA0615hypothetical proteinPA0616hypothetical proteinPA0617probable bacteriophage proteinPA0618probable bacteriophage proteinPA0619probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0639hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0632probable bacteriophage proteinPA0641probable bacteriophage proteinPA0625hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0632probable bacteri  | PA0352 | probable transporter                                |
| PA0459probable ClpA/B protease ATP binding subunitPA0492conserved hypothetical proteinPA0493probable biotin-requiring enzymePA0494probable acyl-CoA carboxylase subunitPA0495hypothetical proteinPA0555fructose-1,6-bisphosphate aldolasePA0586conserved hypothetical proteinPA0587conserved hypothetical proteinPA0613hypothetical proteinPA0614hypothetical proteinPA0615hypothetical proteinPA0616hypothetical proteinPA0617probable bacteriophage proteinPA0618probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0618probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0632probable bacteriophage proteinPA0633probable bacteriophage proteinPA0644hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage protein <td>PA0382</td> <td>DNA mismatch repair protein MicA</td>                              | PA0382 | DNA mismatch repair protein MicA                    |
| PA0492conserved hypothetical proteinPA0493probable biotin-requiring enzymePA0494probable acyl-CoA carboxylase subunitPA0495hypothetical proteinPA0495hypothetical proteinPA0555fructose-1,6-bisphosphate aldolasePA0582dihydroneopterin aldolasePA0586conserved hypothetical proteinPA0587conserved hypothetical proteinPA0613hypothetical proteinPA0614hypothetical proteinPA0615hypothetical proteinPA0616hypothetical proteinPA0617probable bacteriophage proteinPA0618probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0632probable bacteriophage proteinPA0633hypothetical proteinPA0634hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0634hypothetical proteinPA0635hypothetical protein   | PA0385 |   |
| PA0493probable biotin-requiring enzymePA0494probable acyl-CoA carboxylase subunitPA0495hypothetical proteinPA0555fructose-1,6-bisphosphate aldolasePA0552dihydroneopterin aldolasePA0586conserved hypothetical proteinPA0587conserved hypothetical proteinPA0613hypothetical proteinPA0614hypothetical proteinPA0615hypothetical proteinPA0616hypothetical proteinPA0617probable bacteriophage proteinPA0618probable bacteriophage proteinPA0619probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0634hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0639proteinPA0631hypothetical proteinPA0631hypothetical proteinPA0631hypothetical proteinPA0634hypothetical proteinPA0635 <td></td> <td></td>  |        |   |
| PA0494probable acyl-CoA carboxylase subunitPA0495hypothetical proteinPA0555fructose-1,6-bisphosphate aldolasePA0582dihydroneopterin aldolasePA0584conserved hypothetical proteinPA0585conserved hypothetical proteinPA0586conserved hypothetical proteinPA0517conserved hypothetical proteinPA0613hypothetical proteinPA0614hypothetical proteinPA0615hypothetical proteinPA0616hypothetical proteinPA0617probable bacteriophage proteinPA0618probable bacteriophage proteinPA0619probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0634hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothe  |        |   |
| PA0495hypothetical proteinPA0555fructose-1,6-bisphosphate aldolasePA0582dihydroneopterin aldolasePA0584conserved hypothetical proteinPA0587conserved hypothetical proteinPA0612repressor, PtrBPA0613hypothetical proteinPA0614hypothetical proteinPA0615hypothetical proteinPA0616hypothetical proteinPA0617probable bacteriophage proteinPA0618probable bacteriophage proteinPA0619probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0634hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0639conserved hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0632probable bacteriophage proteinPA0634hypothetical proteinPA0635hypothetical proteinPA0636   |        |   |
| PA0555fructose-1,6-bisphosphate aldolasePA0582dihydroneopterin aldolasePA0586conserved hypothetical proteinPA0587conserved hypothetical proteinPA0612repressor, PtrBPA0613hypothetical proteinPA0614hypothetical proteinPA0615hypothetical proteinPA0616hypothetical proteinPA0617probable bacteriophage proteinPA0618probable bacteriophage proteinPA0619probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0629conserved hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0632probable bacteriophagePA0633probable ical proteinPA0634hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0639conserved hypothetical protein<   |        |   |
| PA0582dihydroneopterin aldolasePA0586conserved hypothetical proteinPA0587conserved hypothetical proteinPA0612repressor, PtrBPA0613hypothetical proteinPA0614hypothetical proteinPA0615hypothetical proteinPA0616hypothetical proteinPA0617probable bacteriophage proteinPA0618probable bacteriophage proteinPA0619probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0629conserved hypothetical proteinPA0621probable bacteriophage proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0631hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0639conserved hypothetical proteinPA0630hypothetical proteinPA0636hypothetical proteinPA063  |        |   |
| PA0586conserved hypothetical proteinPA0587conserved hypothetical proteinPA0612repressor, PtrBPA0613hypothetical proteinPA0614hypothetical proteinPA0615hypothetical proteinPA0616hypothetical proteinPA0617probable bacteriophage proteinPA0618probable bacteriophage proteinPA0619probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0629conserved hypothetical proteinPA0620probable bacteriophage proteinPA0621probable bacteriophage proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0638conserved hypothetical proteinPA0631hypothetical proteinPA0633hypothetical proteinPA0634hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0639conserved hypothetical proteinPA0636hypothetical protein <td></td> <td></td>  |        |   |
| PA0587conserved hypothetical proteinPA0612repressor, PtrBPA0613hypothetical proteinPA0614hypothetical proteinPA0615hypothetical proteinPA0616hypothetical proteinPA0617probable bacteriophage proteinPA0618probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0629probable bacteriophage proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0633hypothetical proteinPA0634hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0639conserved hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0632probable bacteriophage proteinPA0634hypothetical proteinPA0635hypothetical proteinPA0636hypot   |        |   |
| PA0612repressor, PtrBPA0613hypothetical proteinPA0614hypothetical proteinPA0615hypothetical proteinPA0616hypothetical proteinPA0617probable bacteriophage proteinPA0618probable bacteriophage proteinPA0619probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0629conserved hypothetical proteinPA0629conserved hypothetical proteinPA0629conserved hypothetical proteinPA0629conserved hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0639conserved hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0634hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage protein  |        |   |
| PA0614hypothetical proteinPA0615hypothetical proteinPA0616hypothetical proteinPA0617probable bacteriophage proteinPA0618probable bacteriophage proteinPA0619probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0629conserved hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0633hypothetical proteinPA0634hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0639conserved hypothetical proteinPA0639conserved hypothetical proteinPA0640probable bacteriophage proteinPA0641probable bacteriophage proteinPA0642hypothetical protein   | PA0612 |   |
| PA0615hypothetical proteinPA0616hypothetical proteinPA0617probable bacteriophage proteinPA0618probable bacteriophage proteinPA0619probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0629conserved hypothetical proteinPA0629conserved hypothetical proteinPA0629proserved hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0639conserved hypothetical proteinPA0630hypothetical proteinPA0634hypothetical proteinPA0635hypothetical proteinPA0639conserved hypothetical proteinPA0639conserved hypothetical proteinPA0640probable bacteriophage proteinPA0641probable bacteriophage proteinPA0642hypothetical protein   | PA0613 | hypothetical protein                                |
| PA0616hypothetical proteinPA0617probable bacteriophage proteinPA0618probable bacteriophage proteinPA0619probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0629conserved hypothetical proteinPA0628conserved hypothetical proteinPA0629conserved hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0639conserved hypothetical proteinPA0630hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0639conserved hypothetical proteinPA0640probable bacteriophage proteinPA0641probable bacteriophage protein   | PA0614 | hypothetical protein                                |
| PA0617probable bacteriophage proteinPA0618probable bacteriophage proteinPA0619probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0629conserved hypothetical proteinPA0629conserved hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0639conserved hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0640probable bacteriophage proteinPA0641probable bacteriophage proteinPA0642hypothetical protein   | PA0615 |   |
| PA0618probable bacteriophage proteinPA0619probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0629conserved hypothetical proteinPA0629conserved hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0640probable bacteriophage proteinPA0641probable bacteriophage proteinPA0642hypothetical protein   |        |   |
| PA0619probable bacteriophage proteinPA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0629conserved hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0639conserved hypothetical proteinPA0640probable bacteriophage proteinPA0641probable bacteriophage proteinPA0642hypothetical protein   |        |   |
| PA0620probable bacteriophage proteinPA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0629conserved hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0639conserved hypothetical proteinPA0636hypothetical proteinPA0637probable bacteriophage proteinPA0640probable bacteriophage proteinPA0640probable bacteriophage proteinPA0641probable bacteriophage proteinPA0642hypothetical protein   |        |   |
| PA0621conserved hypothetical proteinPA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0629conserved hypothetical proteinPA0629conserved hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0633hypothetical proteinPA0634hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0640probable bacteriophage proteinPA0641probable bacteriophage proteinPA0642hypothetical protein   |        |   |
| PA0622probable bacteriophage proteinPA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0629conserved hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0639conserved hypothetical proteinPA0634hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0640probable bacteriophage proteinPA0641probable bacteriophage proteinPA0642hypothetical protein   |        |   |
| PA0623probable bacteriophage proteinPA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0629conserved hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0639conserved hypothetical proteinPA0634probable bacteriophage proteinPA0637probable bacteriophage proteinPA0638probable bacteriophage proteinPA0640probable bacteriophage proteinPA0641probable bacteriophage proteinPA0642hypothetical protein   |        |   |
| PA0624hypothetical proteinPA0625hypothetical proteinPA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0629conserved hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0639conserved hypothetical proteinPA0634probable bacteriophage proteinPA0637probable bacteriophage proteinPA0638probable bacteriophage proteinPA0640probable bacteriophage proteinPA0641probable bacteriophage protein   |        |   |
| PA0626hypothetical proteinPA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0629conserved hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0634hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0639conserved hypothetical proteinPA0640probable bacteriophage proteinPA0641probable bacteriophage proteinPA0642hypothetical protein   |        |   |
| PA0627conserved hypothetical proteinPA0628conserved hypothetical proteinPA0629conserved hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0634hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0639conserved hypothetical proteinPA0640probable bacteriophage proteinPA0641probable bacteriophage protein   |        |   |
| PA0628conserved hypothetical proteinPA0629conserved hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0634hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0640probable bacteriophage proteinPA0641probable bacteriophage proteinPA0642hypothetical protein   | PA0626 | hypothetical protein                                |
| PA0629conserved hypothetical proteinPA0630hypothetical proteinPA0631hypothetical proteinPA0634hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0640probable bacteriophage proteinPA0641probable bacteriophage proteinPA0642hypothetical protein   | PA0627 |   |
| PA0630hypothetical proteinPA0631hypothetical proteinPA0634hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0640probable bacteriophage proteinPA0641probable bacteriophage proteinPA0642hypothetical protein   |        |   |
| PA0631hypothetical proteinPA0634hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0639conserved hypothetical proteinPA0640probable bacteriophage proteinPA0641probable bacteriophage proteinPA0642hypothetical protein   |        |   |
| PA0634hypothetical proteinPA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0639conserved hypothetical proteinPA0640probable bacteriophage proteinPA0641probable bacteriophage proteinPA0642hypothetical protein   |        |   |
| PA0635hypothetical proteinPA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0639conserved hypothetical proteinPA0640probable bacteriophage proteinPA0641probable bacteriophage proteinPA0642hypothetical protein   |        |   |
| PA0636hypothetical proteinPA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0639conserved hypothetical proteinPA0640probable bacteriophage proteinPA0641probable bacteriophage proteinPA0642hypothetical protein   |        |   |
| PA0637conserved hypothetical proteinPA0638probable bacteriophage proteinPA0639conserved hypothetical proteinPA0640probable bacteriophage proteinPA0641probable bacteriophage proteinPA0642hypothetical protein   |        |   |
| PA0638probable bacteriophage proteinPA0639conserved hypothetical proteinPA0640probable bacteriophage proteinPA0641probable bacteriophage proteinPA0642hypothetical protein   |        |   |
| PA0639conserved hypothetical proteinPA0640probable bacteriophage proteinPA0641probable bacteriophage proteinPA0642hypothetical protein   |        |   |
| PA0640probable bacteriophage proteinPA0641probable bacteriophage proteinPA0642hypothetical protein   |        | 1 1 2 1   |
| PA0641probable bacteriophage proteinPA0642hypothetical protein   |        |   |
| 51 1   |        |   |
| PA0643 hypothetical protein  | PA0642 | hypothetical protein                                |
|  | PA0643 | hypothetical protein                                |

### Function class

Membrane proteins Hypothetical, unclassified, unknown Energy metabolism Energy metabolism Energy metabolism Membrane proteins Hypothetical, unclassified, unknown Hypothetical, unclassified, unknown Antibiotic resistance and susceptibility Motility & Attachment Adaptation, Protection Hypothetical, unclassified, unknown Adaptation, Protection Adaptation, Protection Chemotaxis Chemotaxis Putative enzymes Hypothetical, unclassified, unknown Transport of small molecules Membrane proteins Membrane proteins Transport of small molecules Hypothetical, unclassified, unknown Transport of small molecules Membrane proteins DNA replication, recombination, modification and repair Hypothetical, unclassified, unknown Translation, post-translational modification, degradation Hypothetical, unclassified, unknown Putative enzymes Putative enzymes Hypothetical, unclassified, unknown Central intermediary metabolism Biosynthesis of cofactors, prosthetic groups and carriers Hypothetical, unclassified, unknown Hypothetical, unclassified, unknown Transcriptional regulators Hypothetical, unclassified, unknown Hypothetical, unclassified, unknown Hypothetical, unclassified, unknown Related to phage, transposon, or plasmid Related to phage, transposon, or plasmid

| ORF            | Product name  | Function class  |  |
|----------------|---|---|--|
| PA0645         | hypothetical protein  | Related to phage, transposon, or plasmid                  |  |
| PA0646         | hypothetical protein  | Related to phage, transposon, or plasmid                  |  |
| A0669          | probable DNA polymerase alpha chain                               | Putative enzymes  |  |
| A0670          | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |
| A0671          | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |
| PA0743         | probable 3-hydroxyisobutyrate dehydrogenase                       | Carbon compound catabolism                                |  |
| A0745          | probable enoyl-CoA hydratase/isomerase                            | Putative enzymes  |  |
| A0746          | probable acyl-CoA dehydrogenase                                   | Putative enzymes  |  |
| A0747          | probable aldehyde dehydrogenase                                   | Putative enzymes  |  |
| A0873          | transcriptional regulator PhhR                                    | Transcriptional regulators                                |  |
| A0887          | acetyl-coenzyme A synthetase                                      | Carbon compound catabolism                                |  |
| PA0887.1       | aaa   | Non-coding RNA gene                                       |  |
| A0889          | arginine/ornithine transport protein AotQ                         | Membrane proteins   |  |
| A0905          | RsmA, regulator of secondary metabolites                          | Translation, post-translational modification, degradation |  |
| A0915          | conserved hypothetical protein                                    | Hypothetical, unclassified, unknown                       |  |
| A0915<br>A0916 | conserved hypothetical protein                                    | Hypothetical, unclassified, unknown                       |  |
| A0910<br>A0922 | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |
|                |   |   |  |
| A0975          | probable radical activating enzyme                                | Putative enzymes  |  |
| A0976          | conserved hypothetical protein                                    | Hypothetical, unclassified, unknown                       |  |
| A0983          | conserved hypothetical protein                                    | Hypothetical, unclassified, unknown                       |  |
| A0996          | probable coenzyme A ligase  | Biosynthesis of cofactors, prosthetic groups and carriers |  |
| PA1000         | Quinolone signal response protein                                 | Biosynthesis of cofactors, prosthetic groups and carriers |  |
| A1001          | anthranilate synthase component I                                 | Adaptation, Protection                                    |  |
| A1002          | anthranilate synthase component II                                | Amino acid biosynthesis and metabolism                    |  |
| A1041          | probable outer membrane protein precursor                         | Membrane proteins   |  |
| PA1112         | conserved hypothetical protein                                    | Hypothetical, unclassified, unknown                       |  |
| A1150          | pyocin S2   | Adaptation, Protection                                    |  |
| PA1151         | pyocin S2 immunity protein  | Adaptation, Protection                                    |  |
| PA1172         | cytochrome c-type protein NapC                                    | Energy metabolism   |  |
| A1173          | cytochrome c-type protein NapB precursor                          | Energy metabolism   |  |
| A1174          | periplasmic nitrate reductase protein NapA                        | Energy metabolism   |  |
| A1175          | NapD protein of periplasmic nitrate reductase                     | Energy metabolism   |  |
| A1176          | ferredoxin protein NapF   | Energy metabolism   |  |
| A1177          | periplasmic nitrate reductase protein NapE                        | JapE Energy metabolism                                    |  |
| A1228          | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |
| A1320          | cytochrome o ubiquinol oxidase subunit IV                         | Energy metabolism   |  |
| A1321          | cytochrome o ubiquinol oxidase protein CyoE                       | Energy metabolism   |  |
| PA1372         | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |
| A1432          | autoinducer synthesis protein LasI                                | Adaptation, Protection                                    |  |
| PA1466         | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |
| A1552          | probable cytochrome c   | Energy metabolism   |  |
| PA1553         | probable cytochrome c oxidase subunit                             | Energy metabolism   |  |
| A1554          | probable cytochrome oxidase subunit (cbb3-type)                   | Energy metabolism   |  |
| A1574          | conserved hypothetical protein                                    | Hypothetical, unclassified, unknown                       |  |
| A1581          | succinate dehydrogenase (C subunit)                               | Energy metabolism   |  |
| A1581<br>A1582 | succinate dehydrogenase (D subunit)                               | Energy metabolism   |  |
| A1582<br>A1583 | succinate dehydrogenase (A subunit)                               | Energy metabolism   |  |
| A1585<br>A1584 | succinate dehydrogenase (A subunit)                               | Energy metabolism   |  |
| A1584<br>A1634 | potassium-transporting ATPase, B chain                            | Transport of small molecules                              |  |
| A1634<br>A1687 | spermidine synthase   | Amino acid biosynthesis and metabolism                    |  |
| A1687<br>A1759 |   | Transcriptional regulators                                |  |
|                | probable transcriptional regulator                                |   |  |
| A1760          | probable transcriptional regulator                                | Transcriptional regulators                                |  |
| A1776          | ECF sigma factor SigX   | Transcriptional regulators                                |  |
| A1837          | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |
| A1845          | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |
| A1852          | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |
| A1865          | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |
| A1930          | probable chemotaxis transducer                                    | Adaptation, Protection                                    |  |
| A1946          | binding protein component precursor of ABC ribose transporter     | Transport of small molecules                              |  |
| A1951          | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |
| A2006          | probable major facilitator superfamily (MFS) transporter          | Membrane proteins   |  |
| A2007          | maleylacetoacetate isomerase                                      | Carbon compound catabolism                                |  |
| A2008          | fumarylacetoacetase   | Carbon compound catabolism                                |  |
| A2009          | homogentisate 1,2-dioxygenase                                     | Carbon compound catabolism                                |  |
| PA2012         | methylcrotonyl-CoA carboxylase, alpha-subunit (biotin-containing) | Carbon compound catabolism                                |  |
| PA2288         | hypothetical protein  | Hypothetical, unclassified, unknown                       |  |
|                |   |   |  |
| PA2290         | glucose dehydrogenase   | Carbon compound catabolism                                |  |

| ORF              | Product name   | Function class   |
|------------------|--|--|
| PA2322           | gluconate permease   | Transport of small molecules   |
| PA2338           | probable binding protein component of ABC maltose/mannitol transporter | Transport of small molecules   |
| PA2339           | probable binding-protein-dependent maltose/mannitol transport protein  | Transport of small molecules   |
| PA2340           | probable binding-protein-dependent maltose/mannitol transport protein  | Transport of small molecules   |
| PA2341           | probable ATP-binding component of ABC maltose/mannitol transporter     | Transport of small molecules   |
| PA2342           | mannitol dehydrogenase   | Carbon compound catabolism   |
| PA2344           | fructokinase   | Carbon compound catabolism   |
| PA2453           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA2552           | probable acyl-CoA dehydrogenase  | Putative enzymes   |
| PA2553           | probable acyl-CoA thiolase   | Putative enzymes   |
| PA2555           | probable AMP-binding enzyme  | Putative enzymes   |
| PA2557           | probable AMP-binding enzyme  | Fatty acid and phospholipid metabolism                                     |
| PA2573           | probable chemotaxis transducer   | Adaptation, Protection   |
| PA2579           | hypothetical protein   | Putative enzymes   |
| PA2619           | initiation factor  | Translation, post-translational modification, degradation                  |
| PA2637           | NADH dehydrogenase I chain A   | Energy metabolism  |
| PA2638           | NADH dehydrogenase I chain B   | Energy metabolism  |
| PA2639           | NADH dehydrogenase I chain C,D   | Energy metabolism  |
| PA2640           | NADH dehydrogenase I chain E   | Energy metabolism  |
| PA2641           | NADH dehydrogenase I chain F   | Energy metabolism  |
| PA2642           | NADH dehydrogenase I chain G   | Energy metabolism  |
| PA2643           | NADH dehydrogenase I chain H   | Energy metabolism  |
| PA2644<br>PA2645 | NADH Dehydrogenase I chain I   | Energy metabolism  |
| PA2645<br>PA2646 | NADH dehydrogenase I chain J<br>NADH dehydrogenase I chain K           | Energy metabolism<br>Energy metabolism                                     |
| PA2640<br>PA2647 | NADH dehydrogenase I chain L   | Energy metabolism  |
| PA2648           | NADH dehydrogenase I chain M   | Energy metabolism  |
| PA2649           | NADH dehydrogenase I chain N   | Energy metabolism  |
| PA2653           | probable transporter   | Membrane proteins  |
| PA2667           | conserved hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA2730           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA2746           | hypothetical protein   | Membrane proteins  |
| PA2911           | probable TonB-dependent receptor                                       | Membrane proteins  |
| PA2937           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA3007           | repressor protein LexA   | Adaptation, Protection   |
| PA3008           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA3009           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA3080           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA3147           | probable glycosyl transferase WbpJ                                     | Putative enzymes   |
| PA3149<br>PA3150 | probable glycosyltransferase WbpH<br>LPS biosynthesis protein WbpG     | Putative enzymes<br>Cell wall / LPS / capsule                              |
| PA3181           | 2-keto-3-deoxy-6-phosphogluconate aldolase                             | Central intermediary metabolism  |
| PA3183           | glucose-6-phosphate 1-dehydrogenase                                    | Energy metabolism  |
| PA3186           | Glucose/carbohydrate outer membrane porin OprB precursor               | Transport of small molecules   |
| PA3187           | probable ATP-binding component of ABC transporter                      | Transport of small molecules   |
| PA3188           | probable permease of ABC sugar transporter                             | Transport of small molecules   |
| PA3189           | probable permease of ABC sugar transporter                             | Transport of small molecules   |
| PA3190           | probable binding protein component of ABC sugar transporter            | Transport of small molecules   |
| PA3194           | phosphogluconate dehydratase   | Energy metabolism  |
| PA3237           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA3282           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA3311           | conserved hypothetical protein   | Membrane proteins  |
| PA3404           | probable outer membrane protein precursor                              | Protein secretion/export apparatus   |
| PA3415           | probable dihydrolipoamide acetyltransferase                            | Energy metabolism  |
| PA3417           | probable pyruvate dehydrogenase E1 component, alpha subunit            | Energy metabolism  |
| PA3418           | leucine dehydrogenase  | Amino acid biosynthesis and metabolism                                     |
| PA3441           | probable molybdopterin-binding protein                                 | Transport of small molecules   |
| PA3442           | probable ATP-binding component of ABC transporter                      | Transport of small molecules   |
| PA3443<br>PA3444 | probable permease of ABC transporter                                   | Membrane proteins  |
|                  | conserved hypothetical protein   | Putative enzymes<br>Hypothetical, unclassified, unknown                    |
| PA3445<br>PA3446 | conserved hypothetical protein<br>conserved hypothetical protein       | Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown |
| PA3446<br>PA3449 | conserved hypothetical protein   | Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown |
| PA3449<br>PA3496 | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA3561           | 1-phosphofructokinase  | Central intermediary metabolism  |
| PA3574           | probable transcriptional regulator                                     | Transcriptional regulators   |
| PA3581           | glycerol uptake facilitator protein                                    | Transport of small molecules   |
| PA3582           | glycerol kinase  | Central intermediary metabolism  |
|                  |  |  |

| ORF              | Product name   | Function class   |
|------------------|--|--|
| PA3584           | glycerol-3-phosphate dehydrogenase                               | Central intermediary metabolism  |
| PA3600           | conserved hypothetical protein                                   | Translation, post-translational modification, degradation                  |
| PA3607           | polyamine transport protein PotA                                 | Transport of small molecules   |
| PA3608           | polyamine transport protein Poth                                 | Membrane proteins  |
| PA3609           | polyamine transport protein PotC                                 | Membrane proteins  |
| PA3610           | polyamine transport protein PotD                                 | Transport of small molecules   |
| PA3616           | conserved hypothetical protein                                   | Hypothetical, unclassified, unknown  |
| PA3617           | RecA protein   | DNA replication, recombination, modification and repair                    |
| PA3621           | ferredoxin I   | Energy metabolism  |
| PA3672           | probable ATP-binding component of ABC transporter                | Transport of small molecules   |
| PA3712           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA3713           | spermidine dehydrogenase, SpdH                                   | Hypothetical, unclassified, unknown  |
| PA3723           | probable FMN oxidoreductase                                      | Putative enzymes   |
| PA3753           | conserved hypothetical protein                                   | Transport of small molecules   |
| PA3767           | conserved hypothetical protein                                   | Hypothetical, unclassified, unknown  |
| PA3787           | conserved hypothetical protein                                   | Hypothetical, unclassified, unknown  |
| PA3808           | conserved hypothetical protein                                   | Hypothetical, unclassified, unknown  |
| PA3870           | molybdopterin biosynthetic protein A1                            | Biosynthesis of cofactors, prosthetic groups and carriers                  |
| PA3871           | probable peptidyl-prolyl cis-trans isomerase, PpiC-type          | Translation, post-translational modification, degradation                  |
| PA3872           | respiratory nitrate reductase gamma chain                        | Energy metabolism  |
| PA3873           | respiratory nitrate reductase delta chain                        | Energy metabolism  |
| PA3874           | respiratory nitrate reductase beta chain                         | Energy metabolism  |
| PA3875           | respiratory nitrate reductase alpha chain                        | Energy metabolism  |
| PA3876           | nitrite extrusion protein 2                                      | Membrane proteins  |
| PA3877           | nitrite extrusion protein 1                                      | Membrane proteins  |
| PA3901           | Fe(III) dicitrate transport protein FecA                         | Membrane proteins  |
| PA3913           | probable protease  | Putative enzymes   |
| PA3922<br>PA3935 | conserved hypothetical protein<br>taurine dioxygenase            | Hypothetical, unclassified, unknown<br>Carbon compound catabolism          |
| PA3967           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA3986           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA4055           | riboflavin synthase alpha chain                                  | Biosynthesis of cofactors, prosthetic groups and carriers                  |
| PA4131           | probable iron-sulfur protein                                     | Putative enzymes   |
| PA4132           | conserved hypothetical protein                                   | Hypothetical, unclassified, unknown  |
| PA4133           | cytochrome c oxidase subunit (cbb3-type)                         | Energy metabolism  |
| PA4141           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA4199           | probable acyl-CoA dehydrogenase                                  | Putative enzymes   |
| PA4236           | catalase   | Adaptation, Protection   |
| PA4390           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA4427           | stringent starvation protein B                                   | Adaptation, Protection   |
| PA4432           | 30S ribosomal protein S9   | Translation, post-translational modification, degradation                  |
| PA4496           | probable binding protein component of ABC transporter            | Transport of small molecules   |
| PA4553           | type 4 fimbrial biogenesis protein PilX                          | Motility & Attachment  |
| PA4563           | 30S ribosomal protein S20  | Central intermediary metabolism  |
| PA4573           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA4582           | conserved hypothetical protein                                   | Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown |
| PA4583<br>PA4584 | conserved hypothetical protein<br>conserved hypothetical protein | Hypothetical, unclassified, unknown  |
| PA4585           | RNA 3'-terminal phosphate cyclase                                | Transcription, RNA processing and degradation                              |
| PA4585<br>PA4607 | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA4612           | conserved hypothetical protein                                   | Hypothetical, unclassified, unknown  |
| PA4642           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA4645           | probable purine/pyrimidine phosphoribosyl transferase            | Nucleotide biosynthesis and metabolism                                     |
| PA4648           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA4651           | probable pili assembly chaperone                                 | Motility & Attachment  |
| PA4672           | peptidyl-tRNA hydrolase  | Translation, post-translational modification, degradation                  |
| PA4687           | ferric iron-binding periplasmic protein HitA                     | Transport of small molecules   |
| PA4688           | iron (III)-transport system permease HitB                        | Membrane proteins  |
| PA4702           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA4703           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA4732           | glucose-6-phosphate isomerase                                    | Carbon compound catabolism   |
| PA4746           | conserved hypothetical protein                                   | Hypothetical, unclassified, unknown  |
| PA4747           | secretion protein SecG   | Membrane proteins  |
| PA4748           | triosephosphate isomerase  | Central intermediary metabolism  |
| PA4753           | conserved hypothetical protein                                   | Hypothetical, unclassified, unknown  |
| PA4759           | dihydrodipicolinate reductase                                    | Amino acid biosynthesis and metabolism                                     |
| PA4763           | DNA repair protein RecN  | DNA replication, recombination, modification and repair                    |
| PA4839           | biosynthetic arginine decarboxylase                              | Amino acid biosynthesis and metabolism                                     |

| ORF              | Product name  | Function class  |
|------------------|---|---|
| PA4840           | conserved hypothetical protein                            | Hypothetical, unclassified, unknown                       |
| PA4840<br>PA4843 | probable two-component response regulator                 | Transcriptional regulators                                |
|                  |   |   |
| PA5024           | conserved hypothetical protein                            | Hypothetical, unclassified, unknown                       |
| PA5049           | 50S ribosomal protein L31                                 | Translation, post-translational modification, degradation |
| PA5074           | probable ATP-binding component of ABC transporter         | Transport of small molecules                              |
| PA5167           | probable c4-dicarboxylate-binding protein                 | Membrane proteins   |
| PA5235           | glycerol-3-phosphate transporter                          | Membrane proteins   |
| PA5351           | Rubredoxin 1  | Carbon compound catabolism                                |
| PA5380           | GbdR  | Transcriptional regulators                                |
| PA5401           | hypothetical protein                                      | Hypothetical, unclassified, unknown                       |
| PA5436           | probable biotin carboxylase subunit of a transcarboxylase | Central intermediary metabolism                           |
| PA5446           | hypothetical protein                                      | Hypothetical, unclassified, unknown                       |
| PA5452           | phosphomannose isomerase/GDP-mannose WbpW                 | Cell wall / LPS / capsule                                 |
| PA5453           | GDP-mannose 4,6-dehydratase                               | Cell wall / LPS / capsule                                 |
| PA5465           | hypothetical protein                                      | Hypothetical, unclassified, unknown                       |
| PA5470           | probable peptide chain release factor                     | Translation, post-translational modification, degradation |
| PA5471           | hypothetical protein                                      | Hypothetical, unclassified, unknown                       |
| PA5479           | proton-glutamate symporter                                | Membrane proteins   |

| ORF Product name |
|------------------|
|------------------|

Function class

# РСР

| PA0224 | probable aldolase   | Putative enzymes  |
|--------|---|---|
| PA0246 | probable major facilitator superfamily (MFS) transporter                              | Membrane proteins   |
| PA0263 | secreted protein Hcp  | Secreted Factors (toxins, enzymes, alginate)              |
| PA0355 | protease PfpI   | Translation, post-translational modification, degradation |
| PA0417 | probable chemotaxis protein   | Chemotaxis  |
| PA0425 | RND multidrug efflux membrane fusion protein MexA precursor                           | Transport of small molecules                              |
| PA0426 | RND multidrug efflux transporter MexB   | Transport of small molecules                              |
| PA0427 | Major intrinsic multiple antibiotic resistance efflux outer membrane protein OprM pre | e Antibiotic resistance and susceptibility                |
| PA0445 | probable transposase  | Related to phage, transposon, or plasmid                  |
| PA0526 | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA0579 | 30S ribosomal protein S21   | Translation, post-translational modification, degradation |
| PA0726 | hypothetical protein of bacteriophage Pf1   | Hypothetical, unclassified, unknown                       |
| PA0868 | conserved hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA1178 | PhoP/Q and low Mg2+ inducible outer membrane protein H1 precursor                     | Membrane proteins   |
| PA1198 | conserved hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA1281 | cobalamin (5'-phosphate) synthase   | Biosynthesis of cofactors, prosthetic groups and carriers |
| PA1393 | adenosine 5'-phosphosulfate (APS) kinase  | Amino acid biosynthesis and metabolism                    |
| PA1493 | sulfate-binding protein of ABC transporter  | Transport of small molecules                              |
| PA1588 | succinyl-CoA synthetase beta chain  | Energy metabolism   |
| PA1656 | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA1658 | conserved hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA1660 | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA1664 | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA1780 | assimilatory nitrite reductase small subunit  | Central intermediary metabolism                           |
| PA2062 | probable pyridoxal-phosphate dependent enzyme   | Putative enzymes  |
| PA2562 | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA2658 | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA2669 | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA2906 | probable oxidoreductase   | Putative enzymes  |
| PA2941 | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA2951 | electron transfer flavoprotein alpha-subunit  | Energy metabolism   |
| PA3136 | probable secretion protein  | Transport of small molecules                              |
| PA3130 | probable major facilitator superfamily (MFS) transporter                              | Membrane proteins   |
|        | probable UDP-glucose/GDP-mannose dehydrogenase WbpA                                   | ·   |
| PA3159 |   | Putative enzymes  |
| PA3162 | 30S ribosomal protein S1  | Translation, post-translational modification, degradation |
| PA3266 | cold acclimation protein B  | Transcriptional regulators                                |
| PA3547 | poly(beta-d-mannuronate) lyase precursor AlgL   | Cell wall / LPS / capsule                                 |
| PA3552 | ArnB<br>mahahla Bagistanga Nadulatian Call Division (BND) offlux transporter          | Cell wall / LPS / capsule                                 |
| PA3676 | probable Resistance-Nodulation-Cell Division (RND) efflux transporter                 | Membrane proteins   |
| PA3677 | probable RND efflux membrane fusion protein precursor                                 | Transport of small molecules                              |
| PA3678 | probable transcriptional regulator  | Transcriptional regulators                                |
| PA3679 | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA3719 | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA3720 | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA3721 | probable transcriptional regulator  | Transcriptional regulators                                |
| PA3814 | L-cysteine desulfurase (pyridoxal phosphate-dependent)                                | Amino acid biosynthesis and metabolism                    |
| PA3815 | conserved hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA3819 | conserved hypothetical protein  | Membrane proteins   |
| PA3905 | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA4033 | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA4037 | probable ATP-binding component of ABC transporter                                     | Transport of small molecules                              |
| PA4053 | 6,7-dimethyl-8-ribityllumazine synthase   | Biosynthesis of cofactors, prosthetic groups and carriers |
| PA4178 | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA4344 | probable hydrolase  | Putative enzymes  |
| PA4386 | GroES protein   | Chaperones & heat shock proteins                          |
| PA4443 | ATP sulfurylase small subunit   | Central intermediary metabolism                           |
| PA4568 | 50S ribosomal protein L21   | Translation, post-translational modification, degradation |
| PA4816 | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA4847 | biotin carboxyl carrier protein (BCCP)  | Fatty acid and phospholipid metabolism                    |
| PA5013 | branched-chain amino acid transferase   | Amino acid biosynthesis and metabolism                    |
| PA5157 | probable transcriptional regulator  | Transcriptional regulators                                |
| PA5158 | probable outer membrane protein precursor   | Transport of small molecules                              |
| PA5159 | multidrug resistance protein  | Transport of small molecules                              |
| PA5160 | drug efflux transporter   | Membrane proteins   |
| PA5182 | hypothetical protein  | Membrane proteins   |
|        | -   | -   |

| ORF    | Product name   | Function class                      |
|--------|--|-------------------------------------|
| PA5212 | hypothetical protein   | Hypothetical, unclassified, unknown |
| PA5216 | probable permease of ABC iron transporter                              | Membrane proteins                   |
| PA5253 | alginate regulatory protein AlgP                                       | Transcriptional regulators          |
| PA5312 | probable aldehyde dehydrogenase  | Putative enzymes                    |
| PA5326 | hypothetical protein   | Hypothetical, unclassified, unknown |
| PA5369 | phosphate ABC transporter, periplasmic phosphate-binding protein, PstS | Transport of small molecules        |
| PA5390 | probable peptidic bond hydrolase                                       | Putative enzymes                    |
| PA5481 | hypothetical protein   | Hypothetical, unclassified, unknown |
| PA5570 | 50S ribosomal protein L34  | Central intermediary metabolism     |

### Copper

| PA0149           | probable sigma-70 factor, ECF subfamily   |
|------------------|---|
| PA0150           | probable transmembrane sensor   |
| PA0397           | probable cation efflux system protein   |
| PA0423           | PasP  |
| PA0450           | probable phosphate transporter  |
| PA0802           | hypothetical protein  |
| PA0913           | probable Mg transporter MgtE  |
| PA0918<br>PA0931 | cytochrome b561<br>ferric enterobactin receptor PirA                                      |
| PA1134           | hypothetical protein  |
| PA1286           | probable major facilitator superfamily (MFS) transporter                                  |
| PA1297           | probable metal transporter  |
| PA1301           | probable transmembrane sensor   |
| PA1302           | probable heme utilization protein precursor   |
| PA1363           | probable sigma-70 factor, ECF subfamily   |
| PA1364           | probable transmembrane sensor   |
| PA1365           | probable siderophore receptor   |
| PA1390           | probable glycosyl transferase   |
| PA1435           | RND efflux membrane fusion protein precursor  |
| PA1436<br>PA1437 | probable RND efflux transporter<br>probable two-component response regulator              |
| PA1437<br>PA1438 | probable two-component response regulator   |
| PA1562           | aconitate hydratase 1   |
| PA1909           | hypothetical protein  |
| PA1910           | probable tonB-dependent receptor protein  |
| PA1911           | probable transmembrane sensor   |
| PA1912           | probable sigma-70 factor, ECF subfamily   |
| PA2064           | copper resistance protein B precursor   |
| PA2065           | copper resistance protein A precursor   |
| PA2277           | ArsR protein  |
| PA2278           | ArsB protein  |
| PA2279           | ArsC protein  |
| PA2384<br>PA2390 | hypothetical protein<br>probable ATP-binding/permease fusion ABC transporter              |
| PA2390<br>PA2391 | probable outer membrane protein precursor   |
| PA2395           | PvdO  |
| PA2406           | hypothetical protein  |
| PA2451           | hypothetical protein  |
| PA2466           | Ferrioxamine receptor FoxA  |
| PA2467           | Anti-sigma factor FoxR  |
| PA2476           | thiol:disulfide interchange protein DsbG  |
| PA2477           | probable thiol:disulfide interchange protein  |
| PA2478           | probable thiol:disulfide interchange protein<br>probable two-component response regulator |
| PA2479<br>PA2480 | probable two-component response regulator   |
| PA2520           | RND divalent metal cation efflux transporter CzcA   |
| PA2521           | RND divalent metal cation efflux membrane fusion protein CzcB precursor                   |
| PA2522           | outer membrane protein precursor CzcC   |
| PA2524           | probable two-component sensor   |
| PA2664           | flavohemoprotein  |
| PA2686           | two-component response regulator PfeR   |
| PA2688           | Ferric enterobactin receptor, outer membrane protein PfeA precursor                       |
| PA2807           | hypothetical protein  |
| PA2809           | two-component response regulator, CopR<br>two-component sensor, CopS                      |
| PA2810<br>PA2826 | probable glutathione peroxidase   |
| PA2826<br>PA2953 | electron transfer flavoprotein-ubiquinone oxidoreductase                                  |
| PA3392           | nitrous-oxide reductase precursor   |
| PA3407           | heme acquisition protein HasAp  |
| PA3408           | Haem uptake outer membrane receptor HasR precursor  |
| PA3409           | probable transmembrane sensor   |
| PA3410           | probable sigma-70 factor, ECF subfamily   |
| PA3411           | hypothetical protein  |
| PA3412           | hypothetical protein  |
| PA3521           | probable outer membrane protein precursor   |
|                  |   |

Transcriptional regulators Membrane proteins Transport of small molecules Secreted Factors (toxins, enzymes, alginate) Membrane proteins Membrane proteins Transport of small molecules Energy metabolism Transport of small molecules Hypothetical, unclassified, unknown Membrane proteins Membrane proteins Membrane proteins Membrane proteins Transcriptional regulators Membrane proteins Transport of small molecules Putative enzymes Transport of small molecules Membrane proteins Transcriptional regulators Two-component regulatory systems Energy metabolism Hypothetical, unclassified, unknown Transport of small molecules Membrane proteins Transcriptional regulators Adaptation, Protection Adaptation, Protection Transcriptional regulators Membrane proteins Adaptation, Protection Hypothetical, unclassified, unknown Membrane proteins Membrane proteins Adaptation, Protection Hypothetical, unclassified, unknown Hypothetical, unclassified, unknown Transport of small molecules Membrane proteins Translation, post-translational modification, degradation Membrane proteins Membrane proteins Transcriptional regulators Two-component regulatory systems Membrane proteins Membrane proteins Membrane proteins Two-component regulatory systems Energy metabolism Transcriptional regulators Membrane proteins Hypothetical, unclassified, unknown Transcriptional regulators Two-component regulatory systems Adaptation, Protection Energy metabolism Energy metabolism Transport of small molecules Transport of small molecules Membrane proteins Transcriptional regulators Hypothetical, unclassified, unknown Hypothetical, unclassified, unknown

Membrane proteins

| ORF    | Product name  | Function class  |
|--------|---|---|
|        |   |   |
| PA3522 | probable Resistance-Nodulation-Cell Division (RND) efflux transporter | Membrane proteins   |
| PA3530 | conserved hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA3601 | conserved hypothetical protein  | Translation, post-translational modification, degradation |
| PA3602 | conserved hypothetical protein  | Membrane proteins   |
| PA3749 | probable major facilitator superfamily (MFS) transporter              | Membrane proteins   |
| PA3920 | probable metal transporting P-type ATPase                             | Membrane proteins   |
| PA3921 | probable transcriptional regulator                                    | Transcriptional regulators                                |
| PA4158 | ferric enterobactin transport protein FepC                            | Transport of small molecules                              |
| PA4159 | ferrienterobactin-binding periplasmic protein precursor FepB          | Transport of small molecules                              |
| PA4160 | ferric enterobactin transport protein FepD                            | Membrane proteins   |
| PA4192 | probable ATP-binding component of ABC transporter                     | Transport of small molecules                              |
| PA4194 | probable permease of ABC transporter                                  | Membrane proteins   |
| PA4195 | probable binding protein component of ABC transporter                 | Transport of small molecules                              |
| PA4513 | probable oxidoreductase   | Putative enzymes  |
| PA4514 | probable outer membrane receptor for iron transport                   | Transport of small molecules                              |
| PA4515 | conserved hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA4655 | ferrochelatase  | Biosynthesis of cofactors, prosthetic groups and carriers |
| PA4675 | probable TonB-dependent receptor                                      | Transport of small molecules                              |
| PA4705 | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA4706 | probable ATP-binding component of ABC transporter                     | Transport of small molecules                              |
| PA4707 | probable permease of ABC transporter                                  | Membrane proteins   |
| PA4711 | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA4818 | conserved hypothetical protein  | Membrane proteins   |
| PA4819 | probable glycosyl transferase   | Putative enzymes  |
| PA4820 | hypothetical protein  | Hypothetical, unclassified, unknown                       |
| PA4825 | Mg(2+) transport ATPase, P-type 2                                     | Transport of small molecules                              |
| PA5230 | probable permease of ABC transporter                                  | Membrane proteins   |
| PA5231 | probable ATP-binding/permease fusion ABC transporter                  | Membrane proteins   |
| PA5300 | cytochrome c5   | Energy metabolism   |
| PA5304 | D-amino acid dehydrogenase, small subunit                             | Energy metabolism   |
| PA5531 | TonB protein  | Transport of small molecules                              |

# ORF Product name

### Function class

#### Sulfate

| PA0192 | probable TonB-dependent receptor                         |
|--------|--|
| PA0193 | hypothetical protein                                     |
| PA0194 | hypothetical protein                                     |
| PA0197 | hypothetical protein                                     |
| PA0198 | transport protein ExbB                                   |
| PA0199 | transport protein ExbD                                   |
| PA2085 | probable ring-hydroxylating dioxygenase small subunit    |
| PA2086 | probable epoxide hydrolase                               |
| PA2092 | probable major facilitator superfamily (MFS) transporter |
| PA2292 | hypothetical protein                                     |
| PA2293 | hypothetical protein                                     |
| PA2294 | probable ATP-binding component of ABC transporter        |
| PA2295 | probable permease of ABC transporter                     |
| PA2296 | hypothetical protein                                     |
| PA2297 | probable ferredoxin                                      |
| PA2298 | probable oxidoreductase                                  |
| PA2324 | hypothetical protein                                     |
| PA2325 | hypothetical protein                                     |
| PA2326 | hypothetical protein                                     |
| PA2346 | conserved hypothetical protein                           |
| PA2347 | hypothetical protein                                     |
| PA2348 | conserved hypothetical protein                           |
| PA2349 | conserved hypothetical protein                           |
| PA2350 | probable ATP-binding component of ABC transporter        |
| PA2351 | probable permease of ABC transporter                     |
| PA2354 | probable transcriptional regulator                       |
| PA2355 | probable FMNH2-dependent monooxygenase                   |
| PA2356 | methanesulfonate sulfonatase MsuD                        |
| PA2357 | NADH-dependent FMN reductase MsuE                        |
|        |  |

Membrane proteins Putative enzymes Putative enzymes Transport of small molecules Transport of small molecules Transport of small molecules Carbon compound catabolism Carbon compound catabolism Membrane proteins Hypothetical, unclassified, unknown Hypothetical, unclassified, unknown Transport of small molecules Membrane proteins Hypothetical, unclassified, unknown Energy metabolism Putative enzymes Hypothetical, unclassified, unknown Transport of small molecules Membrane proteins Transcriptional regulators Putative enzymes Carbon compound catabolism Carbon compound catabolism

### ORF Product name

### Function class

#### **Osmotic shock**

| PA0236 | probable transcriptional regulator                   |
|--------|--|
| PA0448 | probable transcriptional regulator                   |
| PA1159 | probable cold-shock protein                          |
| PA1226 | probable transcriptional regulator                   |
| PA1243 | probable sensor/response regulator hybrid            |
| PA1690 | translocation protein in type III secretion          |
| PA1692 | probable translocation protein in type III secretion |
| PA1693 | translocation protein in type III secretion          |
| PA1694 | translocation protein in type III secretion          |
| PA1695 | translocation protein in type III secretion          |
| PA1697 | ATP synthase in type III secretion system            |
| PA1702 | conserved hypothetical protein in type III secretion |
| PA1703 | type III secretory apparatus protein PcrD            |
| PA1704 | transcriptional regulator protein PcrR               |
| PA1724 | type III export protein PscK                         |
| PA1725 | type III export protein PscL                         |
| PA2147 | catalase HPII  |
| PA2152 | probable trehalose synthase                          |
| PA2417 | probable transcriptional regulator                   |
| PA2665 | probable transcriptional regulator                   |
| PA3049 | ribosome modulation factor                           |
| PA3459 | probable glutamine amidotransferase                  |
| PA3462 | probable sensor/response regulator hybrid            |
| PA3689 | probable transcriptional regulator                   |
| PA3831 | leucine aminopeptidase                               |
| PA3843 | hypothetical protein                                 |
| PA4135 | probable transcriptional regulator                   |
| PA4581 | transcriptional regulator RtcR                       |
| PA5483 | two-component response regulator AlgB                |
| PA5484 | probable two-component sensor                        |
|        |  |

Transcriptional regulators Transcriptional regulators Transcriptional regulators Transcriptional regulators Two-component regulatory systems Protein secretion/export apparatus Transcriptional regulators Protein secretion/export apparatus Protein secretion/export apparatus Adaptation, Protection Putative enzymes Transcriptional regulators Transcriptional regulators Translation, post-translational modification, degradation Amino acid biosynthesis and metabolism Two-component regulatory systems Transcriptional regulators Transcription, RNA processing and degradation Hypothetical, unclassified, unknown Transcriptional regulators Transcriptional regulators Transcriptional regulators Two-component regulatory systems

# Function class

# Biofilms

| PA0026   | phospholipase C, PlcB  | Hypothetical, unclassified, unknown   |
|--|--|---|
| PA0027   | hypothetical protein   | Hypothetical, unclassified, unknown   |
| PA0028   | hypothetical protein   | Hypothetical, unclassified, unknown   |
| PA0038   | hypothetical protein   | Hypothetical, unclassified, unknown   |
| PA0040   | conserved hypothetical protein   | Hypothetical, unclassified, unknown   |
| PA0046   | hypothetical protein   | Hypothetical, unclassified, unknown   |
| PA0047   | hypothetical protein   | Hypothetical, unclassified, unknown   |
| PA0061   | hypothetical protein   | Hypothetical, unclassified, unknown   |
| PA0062   | hypothetical protein   | Hypothetical, unclassified, unknown   |
| PA0074   | serine/threonine protein kinase PpkA   | Adaptation, Protection  |
| PA0075   | probable phosphoprotein phosphatase  | Putative enzymes  |
| PA0076   | hypothetical protein   | Hypothetical, unclassified, unknown   |
| PA0083   | conserved hypothetical protein   | Hypothetical, unclassified, unknown   |
| PA0086   | hypothetical protein   | Hypothetical, unclassified, unknown   |
| PA0087   | hypothetical protein   | Hypothetical, unclassified, unknown   |
| PA0088   | hypothetical protein   | Hypothetical, unclassified, unknown   |
| PA0089   | hypothetical protein   | Hypothetical, unclassified, unknown   |
| PA0090   | probable ClpA/B-type chaperone   | Translation, post-translational modification, degradation   |
| PA0090   | conserved hypothetical protein   | Hypothetical, unclassified, unknown   |
| PA0092   | hypothetical protein   | Hypothetical, unclassified, unknown   |
| PA0092<br>PA0094   | hypothetical protein   | Hypothetical, unclassified, unknown   |
| PA0102   |  |   |
|  | probable carbonic anhydrase  | Central intermediary metabolism   |
| PA0126   | hypothetical protein   | Hypothetical, unclassified, unknown   |
| PA0134   | probable guanine deaminase   | Nucleotide biosynthesis and metabolism  |
| PA0170   | hypothetical protein   | Hypothetical, unclassified, unknown   |
| PA0224   | probable aldolase  | Putative enzymes  |
| PA0237   | probable oxidoreductase  | Putative enzymes  |
| PA0238   | hypothetical protein   | Hypothetical, unclassified, unknown   |
| PA0252   | hypothetical protein   | Hypothetical, unclassified, unknown   |
| PA0276   | hypothetical protein   | Membrane proteins   |
| PA0320   | conserved hypothetical protein   | Hypothetical, unclassified, unknown   |
| PA0327   | hypothetical protein   | Hypothetical, unclassified, unknown   |
| PA0346   | hypothetical protein   | Hypothetical, unclassified, unknown   |
| PA0347   | glycerophosphoryl diester phosphodiesterase, periplasmic   | Fatty acid and phospholipid metabolism  |
| PA0355   | protease PfpI  | Translation, post-translational modification, degradation   |
| PA0376   | sigma factor RpoH  | Transcriptional regulators  |
| PA0409   | twitching motility protein PilH  | Two-component regulatory systems  |
| PA0410   | twitching motility protein Pill  | Chemotaxis  |
| PA0411   | twitching motility protein PilJ  | Chemotaxis  |
| PA0413   | still frameshift probable component of chemotactic signal transduction system  | Two-component regulatory systems  |
| PA0414   | probable methylesterase  | Chemotaxis  |
| PA0430   | 5,10-methylenetetrahydrofolate reductase   | Amino acid biosynthesis and metabolism  |
| PA0431   | hypothetical protein   | Hypothetical, unclassified, unknown   |
| PA0432   |  |   |
| FA0452   | S-adenosyl-L-homocysteine hydrolase  | Amino acid biosynthesis and metabolism  |
| PA0432<br>PA0446   | S-adenosyl-L-homocysteine hydrolase<br>conserved hypothetical protein  |   |
|  |  | Amino acid biosynthesis and metabolism  |
| PA0446   | conserved hypothetical protein   | Amino acid biosynthesis and metabolism<br>Hypothetical, unclassified, unknown   |
| PA0446<br>PA0451   | conserved hypothetical protein<br>conserved hypothetical protein   | Amino acid biosynthesis and metabolism<br>Hypothetical, unclassified, unknown<br>Membrane proteins  |
| PA0446<br>PA0451<br>PA0452   | conserved hypothetical protein<br>conserved hypothetical protein<br>probable stomatin-like protein   | Amino acid biosynthesis and metabolism<br>Hypothetical, unclassified, unknown<br>Membrane proteins<br>Membrane proteins<br>Hypothetical, unclassified, unknown  |
| PA0446<br>PA0451<br>PA0452<br>PA0462   | conserved hypothetical protein<br>conserved hypothetical protein<br>probable stomatin-like protein<br>hypothetical protein<br>two-component sensor CreC  | Amino acid biosynthesis and metabolism<br>Hypothetical, unclassified, unknown<br>Membrane proteins<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Two-component regulatory systems  |
| PA0446<br>PA0451<br>PA0452<br>PA0462<br>PA0464<br>PA0465   | conserved hypothetical protein<br>conserved hypothetical protein<br>probable stomatin-like protein<br>hypothetical protein<br>two-component sensor CreC<br>inner membrane protein CreD   | Amino acid biosynthesis and metabolism<br>Hypothetical, unclassified, unknown<br>Membrane proteins<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Two-component regulatory systems<br>Membrane proteins   |
| PA0446<br>PA0451<br>PA0452<br>PA0462<br>PA0464<br>PA0465<br>PA0466   | conserved hypothetical protein<br>conserved hypothetical protein<br>probable stomatin-like protein<br>hypothetical protein<br>two-component sensor CreC<br>inner membrane protein CreD<br>hypothetical protein   | Amino acid biosynthesis and metabolism<br>Hypothetical, unclassified, unknown<br>Membrane proteins<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Two-component regulatory systems<br>Membrane proteins<br>Hypothetical, unclassified, unknown  |
| PA0446<br>PA0451<br>PA0452<br>PA0462<br>PA0464<br>PA0465<br>PA0466<br>PA0490   | conserved hypothetical protein<br>conserved hypothetical protein<br>probable stomatin-like protein<br>hypothetical protein<br>two-component sensor CreC<br>inner membrane protein CreD<br>hypothetical protein<br>hypothetical protein   | Amino acid biosynthesis and metabolism<br>Hypothetical, unclassified, unknown<br>Membrane proteins<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Two-component regulatory systems<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown   |
| PA0446<br>PA0451<br>PA0452<br>PA0462<br>PA0464<br>PA0465<br>PA0466<br>PA0490<br>PA0515   | conserved hypothetical protein<br>conserved hypothetical protein<br>probable stomatin-like protein<br>hypothetical protein<br>two-component sensor CreC<br>inner membrane protein CreD<br>hypothetical protein<br>hypothetical protein<br>probable transcriptional regulator   | Amino acid biosynthesis and metabolism<br>Hypothetical, unclassified, unknown<br>Membrane proteins<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Two-component regulatory systems<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Biosynthesis of cofactors, prosthetic groups and carriers  |
| PA0446<br>PA0451<br>PA0452<br>PA0462<br>PA0464<br>PA0465<br>PA0466<br>PA0490<br>PA0515<br>PA0516   | conserved hypothetical protein<br>conserved hypothetical protein<br>probable stomatin-like protein<br>hypothetical protein<br>two-component sensor CreC<br>inner membrane protein CreD<br>hypothetical protein<br>hypothetical protein<br>probable transcriptional regulator<br>heme d1 biosynthesis protein NirF  | Amino acid biosynthesis and metabolism<br>Hypothetical, unclassified, unknown<br>Membrane proteins<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Two-component regulatory systems<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Biosynthesis of cofactors, prosthetic groups and carriers<br>Energy metabolism   |
| PA0446<br>PA0451<br>PA0452<br>PA0462<br>PA0464<br>PA0465<br>PA0466<br>PA0466<br>PA0490<br>PA0515<br>PA0516<br>PA0523   | conserved hypothetical protein<br>conserved hypothetical protein<br>probable stomatin-like protein<br>hypothetical protein<br>two-component sensor CreC<br>inner membrane protein CreD<br>hypothetical protein<br>probable transcriptional regulator<br>heme d1 biosynthesis protein NirF<br>nitric-oxide reductase subunit C  | Amino acid biosynthesis and metabolism<br>Hypothetical, unclassified, unknown<br>Membrane proteins<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Two-component regulatory systems<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Biosynthesis of cofactors, prosthetic groups and carriers<br>Energy metabolism<br>Energy metabolism  |
| PA0446<br>PA0451<br>PA0452<br>PA0462<br>PA0464<br>PA0465<br>PA0466<br>PA0490<br>PA0515<br>PA0516<br>PA0523<br>PA0526   | conserved hypothetical protein<br>conserved hypothetical protein<br>probable stomatin-like protein<br>hypothetical protein<br>two-component sensor CreC<br>inner membrane protein CreD<br>hypothetical protein<br>probable transcriptional regulator<br>heme d1 biosynthesis protein NirF<br>nitric-oxide reductase subunit C<br>hypothetical protein  | Amino acid biosynthesis and metabolism<br>Hypothetical, unclassified, unknown<br>Membrane proteins<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Two-component regulatory systems<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Biosynthesis of cofactors, prosthetic groups and carriers<br>Energy metabolism<br>Energy metabolism<br>Hypothetical, unclassified, unknown   |
| PA0446<br>PA0451<br>PA0452<br>PA0462<br>PA0464<br>PA0465<br>PA0466<br>PA0490<br>PA0515<br>PA0516<br>PA0523<br>PA0526<br>PA0529   | conserved hypothetical protein<br>conserved hypothetical protein<br>probable stomatin-like protein<br>hypothetical protein<br>two-component sensor CreC<br>inner membrane protein CreD<br>hypothetical protein<br>hypothetical protein<br>probable transcriptional regulator<br>heme d1 biosynthesis protein NirF<br>nitric-oxide reductase subunit C<br>hypothetical protein<br>conserved hypothetical protein  | Amino acid biosynthesis and metabolism<br>Hypothetical, unclassified, unknown<br>Membrane proteins<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Two-component regulatory systems<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Biosynthesis of cofactors, prosthetic groups and carriers<br>Energy metabolism<br>Energy metabolism<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown  |
| PA0446<br>PA0451<br>PA0452<br>PA0462<br>PA0464<br>PA0465<br>PA0466<br>PA0466<br>PA0490<br>PA0515<br>PA0516<br>PA0523<br>PA0526<br>PA0529<br>PA0530   | conserved hypothetical protein<br>conserved hypothetical protein<br>probable stomatin-like protein<br>hypothetical protein<br>two-component sensor CreC<br>inner membrane protein CreD<br>hypothetical protein<br>hypothetical protein<br>probable transcriptional regulator<br>heme d1 biosynthesis protein NirF<br>nitric-oxide reductase subunit C<br>hypothetical protein<br>conserved hypothetical protein<br>probable class III pyridoxal phosphate-dependent aminotransferase   | Amino acid biosynthesis and metabolism<br>Hypothetical, unclassified, unknown<br>Membrane proteins<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Two-component regulatory systems<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Biosynthesis of cofactors, prosthetic groups and carriers<br>Energy metabolism<br>Energy metabolism<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Putative enzymes            |
| PA0446<br>PA0451<br>PA0452<br>PA0462<br>PA0464<br>PA0465<br>PA0466<br>PA0490<br>PA0515<br>PA0516<br>PA0523<br>PA0526<br>PA0529<br>PA0530<br>PA0531   | conserved hypothetical protein<br>conserved hypothetical protein<br>probable stomatin-like protein<br>hypothetical protein<br>two-component sensor CreC<br>inner membrane protein CreD<br>hypothetical protein<br>probable transcriptional regulator<br>heme d1 biosynthesis protein NirF<br>nitrie-oxide reductase subunit C<br>hypothetical protein<br>conserved hypothetical protein<br>probable class III pyridoxal phosphate-dependent aminotransferase<br>probable glutamine amidotransferase  | Amino acid biosynthesis and metabolism<br>Hypothetical, unclassified, unknown<br>Membrane proteins<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Two-component regulatory systems<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Biosynthesis of cofactors, prosthetic groups and carriers<br>Energy metabolism<br>Energy metabolism<br>Energy metabolism<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Putative enzymes<br>Putative enzymes  |
| PA0446<br>PA0451<br>PA0452<br>PA0462<br>PA0464<br>PA0465<br>PA0466<br>PA0490<br>PA0515<br>PA0516<br>PA0523<br>PA0523<br>PA0526<br>PA0529<br>PA0530<br>PA0531<br>PA0543                               | conserved hypothetical protein<br>conserved hypothetical protein<br>probable stomatin-like protein<br>hypothetical protein<br>two-component sensor CreC<br>inner membrane protein CreD<br>hypothetical protein<br>hypothetical protein<br>probable transcriptional regulator<br>heme d1 biosynthesis protein NirF<br>nitrie-oxide reductase subunit C<br>hypothetical protein<br>conserved hypothetical protein<br>probable class III pyridoxal phosphate-dependent aminotransferase<br>probable glutamine amidotransferase<br>hypothetical protein  | Amino acid biosynthesis and metabolism<br>Hypothetical, unclassified, unknown<br>Membrane proteins<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Two-component regulatory systems<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Biosynthesis of cofactors, prosthetic groups and carriers<br>Energy metabolism<br>Energy metabolism<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Putative enzymes<br>Putative enzymes<br>Hypothetical, unclassified, unknown  |
| PA0446<br>PA0451<br>PA0452<br>PA0462<br>PA0464<br>PA0465<br>PA0466<br>PA0490<br>PA0515<br>PA0516<br>PA0523<br>PA0523<br>PA0526<br>PA0529<br>PA0530<br>PA0531<br>PA0543<br>PA0547                     | conserved hypothetical protein<br>conserved hypothetical protein<br>probable stomatin-like protein<br>hypothetical protein<br>two-component sensor CreC<br>inner membrane protein CreD<br>hypothetical protein<br>hypothetical protein<br>probable transcriptional regulator<br>heme d1 biosynthesis protein NirF<br>nitrie-oxide reductase subunit C<br>hypothetical protein<br>conserved hypothetical protein<br>probable class III pyridoxal phosphate-dependent aminotransferase<br>probable glutamine amidotransferase<br>hypothetical protein<br>probable glutamine amidotransferase | Amino acid biosynthesis and metabolism<br>Hypothetical, unclassified, unknown<br>Membrane proteins<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Two-component regulatory systems<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Biosynthesis of cofactors, prosthetic groups and carriers<br>Energy metabolism<br>Energy metabolism<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Putative enzymes<br>Putative enzymes<br>Hypothetical, unclassified, unknown<br>Transcriptional regulators  |
| PA0446<br>PA0451<br>PA0452<br>PA0462<br>PA0466<br>PA0465<br>PA0466<br>PA0490<br>PA0515<br>PA0516<br>PA0523<br>PA0526<br>PA0529<br>PA0529<br>PA0530<br>PA0531<br>PA0543<br>PA0547<br>PA0557           | conserved hypothetical protein<br>conserved hypothetical protein<br>probable stomatin-like protein<br>hypothetical protein<br>two-component sensor CreC<br>inner membrane protein CreD<br>hypothetical protein<br>hypothetical protein<br>probable transcriptional regulator<br>heme d1 biosynthesis protein NirF<br>nitrie-oxide reductase subunit C<br>hypothetical protein<br>conserved hypothetical protein<br>probable class III pyridoxal phosphate-dependent aminotransferase<br>probable glutamine amidotransferase<br>hypothetical protein<br>probable transcriptional regulator  | Amino acid biosynthesis and metabolism<br>Hypothetical, unclassified, unknown<br>Membrane proteins<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Two-component regulatory systems<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Biosynthesis of cofactors, prosthetic groups and carriers<br>Energy metabolism<br>Energy metabolism<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Putative enzymes<br>Putative enzymes<br>Hypothetical, unclassified, unknown<br>Transcriptional regulators<br>Hypothetical, unclassified, unknown |
| PA0446<br>PA0451<br>PA0452<br>PA0462<br>PA0464<br>PA0465<br>PA0466<br>PA0490<br>PA0515<br>PA0516<br>PA0523<br>PA0526<br>PA0529<br>PA0529<br>PA0530<br>PA0531<br>PA0543<br>PA0547<br>PA0557<br>PA0585 | conserved hypothetical protein<br>conserved hypothetical protein<br>probable stomatin-like protein<br>hypothetical protein<br>two-component sensor CreC<br>inner membrane protein CreD<br>hypothetical protein<br>probable transcriptional regulator<br>heme d1 biosynthesis protein NirF<br>nitric-oxide reductase subunit C<br>hypothetical protein<br>conserved hypothetical protein<br>probable class III pyridoxal phosphate-dependent aminotransferase<br>probable glutamine amidotransferase<br>hypothetical protein<br>probable transcriptional regulator                          | Amino acid biosynthesis and metabolism<br>Hypothetical, unclassified, unknown<br>Membrane proteins<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Two-component regulatory systems<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Biosynthesis of cofactors, prosthetic groups and carriers<br>Energy metabolism<br>Energy metabolism<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Putative enzymes<br>Putative enzymes<br>Hypothetical, unclassified, unknown<br>Transcriptional regulators<br>Hypothetical, unclassified, unknown |
| PA0446<br>PA0451<br>PA0452<br>PA0462<br>PA0466<br>PA0465<br>PA0466<br>PA0490<br>PA0515<br>PA0516<br>PA0523<br>PA0526<br>PA0529<br>PA0529<br>PA0530<br>PA0531<br>PA0543<br>PA0547<br>PA0557           | conserved hypothetical protein<br>conserved hypothetical protein<br>probable stomatin-like protein<br>hypothetical protein<br>two-component sensor CreC<br>inner membrane protein CreD<br>hypothetical protein<br>hypothetical protein<br>probable transcriptional regulator<br>heme d1 biosynthesis protein NirF<br>nitrie-oxide reductase subunit C<br>hypothetical protein<br>conserved hypothetical protein<br>probable class III pyridoxal phosphate-dependent aminotransferase<br>probable glutamine amidotransferase<br>hypothetical protein<br>probable transcriptional regulator  | Amino acid biosynthesis and metabolism<br>Hypothetical, unclassified, unknown<br>Membrane proteins<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Two-component regulatory systems<br>Membrane proteins<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Biosynthesis of cofactors, prosthetic groups and carriers<br>Energy metabolism<br>Energy metabolism<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown<br>Putative enzymes<br>Putative enzymes<br>Hypothetical, unclassified, unknown<br>Transcriptional regulators<br>Hypothetical, unclassified, unknown |

| ORF            | Product name   | Function class  |
|----------------|--|---|
| PA0603         | probable ATP-binding component of ABC transporter                              | Transport of small molecules                              |
| PA0674         | hypothetical protein   | Hypothetical, unclassified, unknown                       |
| PA0675         | probable sigma-70 factor, ECF subfamily  | Transcriptional regulators                                |
| A0676          | probable transmembrane sensor  | Membrane proteins   |
| A0693          | transport protein ExbB2  | Membrane proteins   |
| A0696          | hypothetical protein   | Hypothetical, unclassified, unknown                       |
| A0697          | hypothetical protein   | Hypothetical, unclassified, unknown                       |
| A0698          | hypothetical protein   | Hypothetical, unclassified, unknown                       |
| A0699          |  |   |
| PA0707         | probable peptidyl-prolyl cis-trans isomerase, PpiC-type                        | Translation, post-translational modification, degradation |
| PA0732         | transcriptional regulator ToxR   | Transcriptional regulators                                |
|                | hypothetical protein   | Hypothetical, unclassified, unknown                       |
| PA0737         | hypothetical protein   | Hypothetical, unclassified, unknown                       |
| PA0738         | conserved hypothetical protein   | Membrane proteins   |
| PA0752         | conserved hypothetical protein   | Membrane proteins   |
| PA0753         | hypothetical protein   | Membrane proteins   |
| PA0755         | cis-aconitate porin OpdH   | Membrane proteins   |
| PA0765         | positive regulator for alginate biosynthesis MucC                              | Transcriptional regulators                                |
| A0788          | hypothetical protein   | Hypothetical, unclassified, unknown                       |
| A0842          | probable glycosyl transferase  | Putative enzymes  |
| A0853          | probable oxidoreductase  | Putative enzymes  |
| A0877          | probable transcriptional regulator   | Transcriptional regulators                                |
| PA0878         | hypothetical protein   | Hypothetical, unclassified, unknown                       |
| PA0879         | probable acyl-CoA dehydrogenase  | Putative enzymes  |
| PA0880         | probable ring-cleaving dioxygenase   | Putative enzymes  |
| PA0882         | hypothetical protein   | Putative enzymes  |
| PA0884         | probable C4-dicarboxylate-binding periplasmic protein                          | Transport of small molecules                              |
| PA0886         | probable C4-dicarboxylate transporter  | Transport of small molecules                              |
| PA0921         | hypothetical protein   | Membrane proteins   |
| PA0952         | hypothetical protein   | Hypothetical, unclassified, unknown                       |
| PA0985         | pyocin S5  | Membrane proteins   |
| A1029          | hypothetical protein   | Hypothetical, unclassified, unknown                       |
| PA1065         | conserved hypothetical protein   | Hypothetical, unclassified, unknown                       |
| PA1065         | probable short-chain dehydrogenase   | Putative enzymes  |
| PA1078         |  | -   |
|                | flagellar basal-body rod protein FlgC  | Cell wall / LPS / capsule                                 |
| PA1079         | flagellar basal-body rod modification protein FlgD                             | Cell wall / LPS / capsule                                 |
| PA1080         | flagellar hook protein FlgE  | Cell wall / LPS / capsule                                 |
| PA1081         | flagellar basal-body rod protein FlgF  | Cell wall / LPS / capsule                                 |
| PA1084         | flagellar P-ring protein precursor FlgI  | Cell wall / LPS / capsule                                 |
| PA1111         | hypothetical protein   | Hypothetical, unclassified, unknown                       |
| PA1114         | hypothetical protein   | Hypothetical, unclassified, unknown                       |
| PA1115         | hypothetical protein   | Membrane proteins   |
| PA1118         | hypothetical protein   | Membrane proteins   |
| PA1155         | NrdB, tyrosyl radical-harboring component of class Ia ribonucleotide reductase | Nucleotide biosynthesis and metabolism                    |
| PA1168         | hypothetical protein   | Hypothetical, unclassified, unknown                       |
| PA1178         | PhoP/Q and low Mg2+ inducible outer membrane protein H1 precursor              | Membrane proteins   |
| PA1195         | hypothetical protein   | Hypothetical, unclassified, unknown                       |
| PA1216         | hypothetical protein   | Hypothetical, unclassified, unknown                       |
| PA1217         | probable 2-isopropylmalate synthase  | Amino acid biosynthesis and metabolism                    |
| PA1247         | alkaline protease secretion protein AprE                                       | Secreted Factors (toxins, enzymes, alginate)              |
| PA1285         | probable transcriptional regulator   | Transcriptional regulators                                |
| PA1333         | hypothetical protein   | Hypothetical, unclassified, unknown                       |
| PA1345         | hypothetical protein   | Hypothetical, unclassified, unknown                       |
| PA1404         | hypothetical protein   | Hypothetical, unclassified, unknown                       |
| A1470          | probable short-chain dehydrogenase   | Putative enzymes  |
| A1471          | hypothetical protein   | Hypothetical, unclassified, unknown                       |
| A1514          | conserved hypothetical protein   | Hypothetical, unclassified, unknown                       |
| A1514<br>A1517 | conserved hypothetical protein   | Hypothetical, unclassified, unknown                       |
| A1517<br>A1518 | conserved hypothetical protein   | Hypothetical, unclassified, unknown                       |
|                |  |   |
| PA1523         | xanthine dehydrogenase   | Nucleotide biosynthesis and metabolism                    |
| PA1524         | xanthine dehydrogenase   | Nucleotide biosynthesis and metabolism                    |
| PA1560         | hypothetical protein   | Hypothetical, unclassified, unknown                       |
| PA1600         | probable cytochrome c  | Energy metabolism   |
| PA1601         | probable aldehyde dehydrogenase  | Putative enzymes  |
| PA1602         | probable oxidoreductase  | Carbon compound catabolism                                |
| PA1603         | probable transcriptional regulator   | Transcriptional regulators                                |
| PA1606         | hypothetical protein   | Hypothetical, unclassified, unknown                       |
|                |  |   |
| PA1625         | conserved hypothetical protein   | Membrane proteins   |

| ORF              | Product name  | Function class  |
|------------------|---|---|
| PA1730           | conserved hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA1780           | assimilatory nitrite reductase small subunit  | Central intermediary metabolism   |
| PA1786           | conserved hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA1791           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA1847           | conserved hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA1870           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA1889           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA1931           | probable ferredoxin   | Energy metabolism   |
| PA1932           | probable hydroxylase molybdopterin-containing subunit   | Carbon compound catabolism  |
| PA1933           | probable hydroxylase large subunit  | Carbon compound catabolism  |
| PA2017           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2021           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2022           | probable nucleotide sugar dehydrogenase   | Putative enzymes  |
| PA2024           | probable ring-cleaving dioxygenase  | Putative enzymes  |
| PA2042           | probable transporter (membrane subunit)   | Transport of small molecules  |
| PA2071           | elongation factor G   | Translation, post-translational modification, degradation                     |
| PA2076           | probable transcriptional regulator  | Transcriptional regulators  |
| PA2107           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2136           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2140           | probable metallothionein  | Central intermediary metabolism   |
| PA2149           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2155           | probable phospholipase  | Putative enzymes  |
| PA2167           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2168           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2170           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2174           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2175           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2176           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2178           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2179           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2181           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2186           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2188           | probable alcohol dehydrogenase (Zn-dependent)   | Putative enzymes  |
| PA2243           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2365           | conserved hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2366           | conserved hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2367           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2369           | hypothetical protein  | Membrane proteins   |
| PA2441           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2444           | serine hydroxymethyltransferase   | Amino acid biosynthesis and metabolism  |
| PA2445           | glycine cleavage system protein P2  | Central intermediary metabolism   |
| PA2446           | glycine cleavage system protein H2  | Amino acid biosynthesis and metabolism  |
| PA2459           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2462           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2485           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2486           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2491           | probable oxidoreductase   | Putative enzymes  |
| PA2493           | RND multidrug efflux membrane fusion protein MexE precursor                                     | Transport of small molecules  |
| PA2494           | RND multidrug efflux transporter MexF<br>Multidrug efflux outer membrane protein OprN precursor | Transport of small molecules  |
| PA2495           |   | Membrane proteins   |
| PA2533<br>PA2536 | probable sodium:alanine symporter   | Membrane proteins   |
| PA2536<br>PA2562 | probable phosphatidate cytidylyltransferase<br>hypothetical protein                             | Fatty acid and phospholipid metabolism<br>Hypothetical, unclassified, unknown |
| PA2562<br>PA2569 | hypothetical protein  | Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown    |
| PA2569<br>PA2618 | hypothetical protein  | Translation, post-translational modification, degradation                     |
| PA2618<br>PA2635 | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2635<br>PA2658 | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2658<br>PA2659 | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2039<br>PA2708 | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2708<br>PA2718 | probable transcriptional regulator  | Transcriptional regulators  |
| PA2761           | hypothetical protein  | Membrane proteins   |
| PA2701<br>PA2777 | conserved hypothetical protein  | Membrane proteins   |
| PA2777<br>PA2782 | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2782<br>PA2799 | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2803           | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2803<br>PA2804 | hypothetical protein  | Hypothetical, unclassified, unknown   |
| PA2815           | probable acyl-CoA dehydrogenase   | Putative enzymes  |
| PA2816           | hypothetical protein  | Hypothetical, unclassified, unknown   |
|                  | 5 r · · · · · · · · · · · · · ·   |   |

| ORF              | Product name   | Function class   |
|------------------|--|--|
| PA2827           | conserved hypothetical protein   | Hypothetical, unclassified, unknown  |
| A2880            | hypothetical protein   | Hypothetical, unclassified, unknown  |
| A2881            | probable two-component response regulator  | Transcriptional regulators   |
| A2882            | probable two-component sensor  | Two-component regulatory systems   |
| A2883            | hypothetical protein   | Hypothetical, unclassified, unknown  |
| A2895            | hypothetical protein   | Hypothetical, unclassified, unknown  |
| A2931            | probable transcriptional regulator   | Transcriptional regulators   |
| A2939<br>A2969   | probable aminopeptidase<br>fatty acid biosynthesis protein PlsX  | Secreted Factors (toxins, enzymes, alginate)<br>Fatty acid and phospholipid metabolism |
| A2909<br>A2970   | 50S ribosomal protein L32  | Translation, post-translational modification, degradation                              |
| A3023            | conserved hypothetical protein   | Hypothetical, unclassified, unknown  |
| A3031            | hypothetical protein   | Hypothetical, unclassified, unknown  |
| A3079            | hypothetical protein   | Membrane proteins  |
| A3082            | glycine betaine transmethylase   | Amino acid biosynthesis and metabolism   |
| PA3105           | general secretion pathway protein D  | Protein secretion/export apparatus   |
| PA3121           | 3-isopropylmalate dehydratase large subunit  | Amino acid biosynthesis and metabolism   |
| PA3205           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA3218           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| A3219            | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA3221           | CsaA protein   | Protein secretion/export apparatus   |
| PA3229<br>PA3249 | hypothetical protein<br>probable transcriptional regulator   | Hypothetical, unclassified, unknown<br>Transcriptional regulators                      |
| A3249<br>A3273   | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA3274           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA3280           | Pyrophosphate-specific outer membrane porin OprO precursor   | Transport of small molecules   |
| PA3296           | alkaline phosphatase   | Central intermediary metabolism  |
| PA3309           | conserved hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA3319           | non-hemolytic phospholipase C precursor  | Secreted Factors (toxins, enzymes, alginate)   |
| PA3326           | probable Clp-family ATP-dependent protease   | Translation, post-translational modification, degradation                              |
| PA3327           | probable non-ribosomal peptide synthetase  | Adaptation, Protection   |
| PA3329           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA3330           | probable short chain dehydrogenase   | Putative enzymes   |
| PA3331           | cytochrome P450  | Adaptation, Protection   |
| PA3332           | conserved hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA3333<br>PA3334 | 3-oxoacyl-[acyl-carrier-protein] synthase III<br>probable acyl carrier protein                                     | Fatty acid and phospholipid metabolism<br>Fatty acid and phospholipid metabolism       |
| PA3335           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA3336           | probable major facilitator superfamily (MFS) transporter   | Membrane proteins  |
| PA3337           | ADP-L-glycero-D-mannoheptose 6-epimerase   | Cell wall / LPS / capsule  |
| PA3368           | probable acetyltransferase   | Putative enzymes   |
| PA3376           | probable ATP-binding component of ABC transporter  | Transport of small molecules   |
| PA3377           | conserved hypothetical protein   | Transport of small molecules   |
| PA3380           | conserved hypothetical protein   | Transport of small molecules   |
| PA3381           | probable transcriptional regulator   | Transcriptional regulators   |
| PA3382           | phosphonate transport protein PhnE   | Membrane proteins  |
| PA3383           | binding protein component of ABC phosphonate transporter   | Transport of small molecules   |
| PA3384           | ATP-binding component of ABC phosphonate transporter   | Transport of small molecules   |
| PA3389<br>PA3397 | probable ring-cleaving dioxygenase<br>ferredoxinNADP+ reductase  | Putative enzymes   |
| PA3431           | conserved hypothetical protein   | Biosynthesis of cofactors, prosthetic groups and carriers<br>Membrane proteins         |
| PA3431<br>PA3432 | hypothetical protein   | Membrane proteins  |
| PA3436           | hypothetical protein   | Hypothetical, unclassified, unknown  |
| PA3476           | autoinducer synthesis protein RhlI   | Adaptation, Protection   |
| A3478            | rhamnosyltransferase chain B   | Secreted Factors (toxins, enzymes, alginate)   |
| A3484            | hypothetical protein   | Hypothetical, unclassified, unknown  |
| A3540            | GDP-mannose 6-dehydrogenase AlgD   | Cell wall / LPS / capsule  |
| A3541            | alginate biosynthesis protein Alg8   | Cell wall / LPS / capsule  |
| A3542            | alginate biosynthesis protein Alg44  | Cell wall / LPS / capsule  |
| A3543            | alginate biosynthetic protein AlgK precursor   | Cell wall / LPS / capsule  |
| A3544            | Alginate production outer membrane protein AlgE precursor  | Membrane proteins  |
| A3545            | alginate-c5-mannuronan-epimerase AlgG  | Cell wall / LPS / capsule  |
| PA3546           | alginate biosynthesis protein AlgX   | Cell wall / LPS / capsule  |
| PA3547           | poly(beta-d-mannuronate) lyase precursor AlgL  | Cell wall / LPS / capsule  |
| PA3548           | alginate o-acetyltransferase AlgI  | Cell wall / LPS / capsule  |
| PA3549           | alginate o-acetyltransferase AlgJ  | Cell wall / LPS / capsule  |
| PA3550<br>PA3551 | alginate o-acetyltransferase AlgF<br>phosphomannose isomerase / guanosine 5'-diphospho-D-mannose pyrophosphorylase | Cell wall / LPS / capsule<br>Cell wall / LPS / capsule                                 |
|                  |  |  |

| ORF              | Product name  | Function class   |
|------------------|---|--|
| PA3554           | AmA   | Putative enzymes   |
| PA3555           | ArnD  | Adaptation, Protection   |
| PA3556           | inner membrane L-Ara4N transferase ArnT                         | Adaptation, Protection   |
| PA3557           | AmE   | Adaptation, Protection   |
| PA3559           | probable nucleotide sugar dehydrogenase                         | Putative enzymes   |
| PA3569           | 3-hydroxvisobutyrate dehydrogenase                              | Carbon compound catabolism   |
| PA3577           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA3578           | conserved hypothetical protein                                  | Hypothetical, unclassified, unknown  |
| PA3598           | conserved hypothetical protein                                  | Hypothetical, unclassified, unknown  |
| PA3651           | phosphatidate cytidylyltransferase                              | Fatty acid and phospholipid metabolism                                     |
| PA3655           | elongation factor Ts  | Translation, post-translational modification, degradation                  |
| PA3688           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA3692           | probable outer membrane protein precursor                       | Membrane proteins  |
| PA3724           | elastase LasB   | Secreted Factors (toxins, enzymes, alginate)                               |
| PA3743           | tRNA (guanine-N1)-methyltransferase                             | Transcription, RNA processing and degradation                              |
| PA3744           | 16S rRNA processing protein                                     | Transcription, RNA processing and degradation                              |
| PA3819           | conserved hypothetical protein                                  | Membrane proteins  |
| PA3820           | secretion protein SecF  | Protein secretion/export apparatus   |
| PA3821           | secretion protein Sech  | Membrane proteins  |
| PA3821<br>PA3840 | conserved hypothetical protein                                  | Hypothetical, unclassified, unknown  |
| PA3840<br>PA3890 |   | Membrane proteins  |
| PA3890<br>PA3905 | probable permease of ABC transporter                            | 1  |
| PA3905<br>PA3906 | hypothetical protein<br>hypothetical protein                    | Hypothetical, unclassified, unknown<br>Hypothetical, unclassified, unknown |
| PA3900<br>PA3907 |   |  |
|                  | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA3908           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA3909           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA3910           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA3929           | cyanide insensitive terminal oxidase                            | Energy metabolism  |
| PA3944           | conserved hypothetical protein                                  | Hypothetical, unclassified, unknown  |
| PA3952           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA3957           | probable short-chain dehydrogenase                              | Putative enzymes   |
| PA3964           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA4017           | conserved hypothetical protein                                  | Hypothetical, unclassified, unknown  |
| PA4040           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA4063           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA4110           | beta-lactamase precursor  | Adaptation, Protection   |
| PA4111           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA4112           | probable sensor/response regulator hybrid                       | Two-component regulatory systems   |
| PA4154           | conserved hypothetical protein                                  | Hypothetical, unclassified, unknown  |
| PA4155           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA4190           | probable FAD-dependent monooxygenase                            | Putative enzymes   |
| PA4238           | DNA-directed RNA polymerase alpha chain                         | Transcription, RNA processing and degradation                              |
| PA4243           | secretion protein SecY  | Membrane proteins  |
| PA4244           | 50S ribosomal protein L15                                       | Translation, post-translational modification, degradation                  |
| PA4246           | 30S ribosomal protein S5  | Translation, post-translational modification, degradation                  |
| PA4247           | 50S ribosomal protein L18                                       | Translation, post-translational modification, degradation                  |
| PA4248           | 50S ribosomal protein L6  | Translation, post-translational modification, degradation                  |
| PA4255           | 50S ribosomal protein L29                                       | Translation, post-translational modification, degradation                  |
| PA4261           | 50S ribosomal protein L23                                       | Translation, post-translational modification, degradation                  |
| PA4262           | 50S ribosomal protein L4  | Transcription, RNA processing and degradation                              |
| PA4272           | 50S ribosomal protein L10                                       | Translation, post-translational modification, degradation                  |
| PA4293           | two-component sensor PprA                                       | Two-component regulatory systems   |
| PA4317           | hypothetical protein  | Membrane proteins  |
| PA4344           | probable hydrolase  | Putative enzymes   |
| PA4345           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA4348           | conserved hypothetical protein                                  | Hypothetical, unclassified, unknown  |
| PA4350           | conserved hypothetical protein                                  | Hypothetical, unclassified, unknown  |
| PA4351           | probable acyltransferase  | Fatty acid and phospholipid metabolism                                     |
| PA4354           | conserved hypothetical protein                                  | Hypothetical, unclassified, unknown  |
| PA4355           | probable major facilitator superfamily (MFS) transporter        | Membrane proteins  |
| PA4356           | xenobiotic reductase  | Adaptation, Protection   |
| PA4357           | conserved hypothetical protein                                  | Hypothetical, unclassified, unknown  |
| PA4371           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA4407           | cell division protein FtsZ                                      | Cell division  |
| PA4407           | cell division protein FtsA                                      | Cell division  |
| PA4408<br>PA4409 | cell division protein FtsQ                                      | Cell division  |
| 11111107         |   | Cell wall / LPS / capsule  |
| PA4410           |   |  |
| PA4410<br>PA4411 | D-alanineD-alanine ligase<br>UDP-N-acetylmuramatealanine ligase | Cell wall / LPS / capsule  |

| ORF              | Product name  | Function class   |
|------------------|---|--|
| PA4412           | UDP-N-acetylglucosamineN-acetylmuramyl undecaprenol N-acetylglucosamine trai        | Carbon compound catabolism   |
| PA4413           | cell division protein FtsW  | Cell division  |
| PA4414           | UDP-N-acetylmuramoylalanineD-glutamate ligase                                       | Cell wall / LPS / capsule  |
| PA4415           | phospho-N-acetylmuramoyl-pentapeptide-transferase                                   | Cell wall / LPS / capsule  |
| PA4416           | UDP-N-acetylmuramoylalanyl-D-glutamyl-2, 6-diaminopimelateD-alanyl-D-alanyl         |  |
| PA4419           | cell division protein FtsL  | Cell division  |
| PA4421<br>PA4498 | conserved hypothetical protein<br>probable metallopeptidase                         | Hypothetical, unclassified, unknown<br>Translation, post-translational modification, degradation                       |
| PA4498<br>PA4500 | probable binding protein component of ABC transporter                               | Translation, post-translational modification, degradation<br>Transport of small molecules                              |
| PA4502           | probable binding protein component of ABC transporter                               | Transport of small molecules   |
| PA4503           | probable permease of ABC transporter  | Membrane proteins  |
| PA4504           | probable permease of ABC transporter  | Membrane proteins  |
| PA4505           | probable ATP-binding component of ABC transporter                                   | Transport of small molecules   |
| PA4506           | probable ATP-binding component of ABC dipeptide transporter                         | Transport of small molecules   |
| PA4507           | hypothetical protein  | Membrane proteins  |
| PA4525           | type 4 fimbrial precursor PilA  | Motility & Attachment  |
| PA4526           | type 4 fimbrial biogenesis protein PilB   | Motility & Attachment  |
| PA4569           | octaprenyl-diphosphate synthase   | Energy metabolism  |
| PA4575           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA4588<br>PA4589 | glutamate dehydrogenase<br>probable outer membrane protein precursor                | Amino acid biosynthesis and metabolism<br>Membrane proteins  |
| PA4621           | probable oxidoreductase   | Putative enzymes   |
| PA4623           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA4632           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA4657           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA4658           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA4671           | probable ribosomal protein L25  | Adaptation, Protection   |
| PA4674           | conserved hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA4691           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA4697           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA4713<br>PA4742 | hypothetical protein<br>+PNA regulariziding 55 sumthage                             | Hypothetical, unclassified, unknown  |
| PA4742<br>PA4743 | tRNA pseudouridine 55 synthase<br>ribosome-binding factor A                         | Translation, post-translational modification, degradation<br>Translation, post-translational modification, degradation |
| PA4744           | translation initiation factor IF-2  | Translation, post-translational modification, degradation  |
| PA4773           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA4781           | probable two-component response regulator   | Transcriptional regulators   |
| PA4782           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA4785           | probable acyl-CoA thiolase  | Putative enzymes   |
| PA4786           | probable short-chain dehydrogenase  | Putative enzymes   |
| PA4811           | nitrate-inducible formate dehydrogenase, beta subunit                               | Energy metabolism  |
| PA4813           | lipase LipC   | Fatty acid and phospholipid metabolism   |
| PA4844           | probable chemotaxis transducer  | Adaptation, Protection   |
| PA4858<br>PA4862 | conserved hypothetical protein<br>probable ATP-binding component of ABC transporter | Hypothetical, unclassified, unknown<br>Transport of small molecules  |
| PA4870           | conserved hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA4877           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA4879           | conserved hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA4881           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA4896           | probable sigma-70 factor, ECF subfamily   | Transcriptional regulators   |
| PA4933           | hypothetical protein  | Membrane proteins  |
| PA5033           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA5040           | Type 4 fimbrial biogenesis outer membrane protein PilQ precursor                    | Motility & Attachment  |
| PA5041           | type 4 fimbrial biogenesis protein PilP<br>type 4 fimbrial biogenesis protein PilO  | Motility & Attachment  |
| PA5042<br>PA5043 | type 4 fimbrial biogenesis protein PilO<br>type 4 fimbrial biogenesis protein PilN  | Motility & Attachment<br>Motility & Attachment   |
| PA5045<br>PA5044 | type 4 fimbrial biogenesis protein PilM   | Motility & Attachment  |
| PA5075           | probable permease of ABC transporter  | Membrane proteins  |
| PA5097           | probable amino acid permease  | Membrane proteins  |
| PA5099           | probable transporter  | Membrane proteins  |
| PA5116           | probable transcriptional regulator  | Transcriptional regulators   |
| PA5124           | two-component sensor NtrB   | Two-component regulatory systems   |
| PA5125           | two-component response regulator NtrC   | Transcriptional regulators   |
| PA5128           | secretion protein SecB  | Protein secretion/export apparatus   |
| PA5129           | glutaredoxin  | Energy metabolism  |
| PA5139           | hypothetical protein  | Hypothetical, unclassified, unknown  |
| PA5169<br>PA5170 | probable C4-dicarboxylate transporter<br>arginine/ornithine antiporter              | Membrane proteins<br>Membrane proteins   |
| PA5180           | conserved hypothetical protein  | Hypothetical, unclassified, unknown  |
|                  |   |  |

| ORF    | Product name  | Function class                               |
|--------|---|--|
|        |   |  |
| PA5182 | hypothetical protein                                    | Membrane proteins                            |
| PA5183 | hypothetical protein                                    | Membrane proteins                            |
| PA5192 | phosphoenolpyruvate carboxykinase                       | Carbon compound catabolism                   |
| PA5202 | hypothetical protein                                    | Hypothetical, unclassified, unknown          |
| PA5209 | hypothetical protein                                    | Hypothetical, unclassified, unknown          |
| PA5212 | hypothetical protein                                    | Hypothetical, unclassified, unknown          |
| PA5220 | hypothetical protein                                    | Hypothetical, unclassified, unknown          |
| PA5232 | conserved hypothetical protein                          | Hypothetical, unclassified, unknown          |
| PA5262 | alginate biosynthesis protein AlgZ/FimS                 | Secreted Factors (toxins, enzymes, alginate) |
| PA5275 | conserved hypothetical protein                          | Hypothetical, unclassified, unknown          |
| PA5287 | ammonium transporter AmtB                               | Membrane proteins                            |
| PA5299 | hypothetical protein                                    | Hypothetical, unclassified, unknown          |
| PA5322 | phosphomannomutase AlgC                                 | Cell wall / LPS / capsule                    |
| PA5360 | two-component response regulator PhoB                   | Transcriptional regulators                   |
| PA5361 | two-component sensor PhoR                               | Two-component regulatory systems             |
| PA5366 | ATP-binding component of ABC phosphate transporter      | Membrane proteins                            |
| PA5367 | membrane protein component of ABC phosphate transporter | Membrane proteins                            |
| PA5368 | membrane protein component of ABC phosphate transporter | Membrane proteins                            |
| PA5415 | serine hydroxymethyltransferase                         | Amino acid biosynthesis and metabolism       |
| PA5422 | hypothetical protein                                    | Hypothetical, unclassified, unknown          |
| PA5441 | hypothetical protein                                    | Hypothetical, unclassified, unknown          |
| PA5460 | hypothetical protein                                    | Hypothetical, unclassified, unknown          |
| PA5473 | conserved hypothetical protein                          | Membrane proteins                            |
| PA5475 | hypothetical protein                                    | Hypothetical, unclassified, unknown          |
| PA5476 | citrate transporter                                     | Membrane proteins                            |
| PA5498 | probable adhesin  | Motility & Attachment                        |
| PA5499 | transcriptional regulator np20                          | Transcriptional regulators                   |
| PA5524 | probable short-chain dehydrogenase                      | Putative enzymes                             |
| PA5525 | probable transcriptional regulator                      | Transcriptional regulators                   |
| PA5526 | hypothetical protein                                    | Hypothetical, unclassified, unknown          |
| PA5549 | glucosaminefructose-6-phosphate aminotransferase        | Central intermediary metabolism              |
| PA5555 | ATP synthase gamma chain                                | Energy metabolism                            |
| PA5556 | ATP synthase alpha chain                                | Energy metabolism                            |
| PA5557 | ATP synthase delta chain                                | Energy metabolism                            |
| PA5558 | ATP synthase B chain                                    | Energy metabolism                            |
| PA5559 | atp synthase C chain                                    | Energy metabolism                            |
| PA5560 | ATP synthase A chain                                    | Energy metabolism                            |
| PA5568 | conserved hypothetical protein                          | Membrane proteins                            |
|        | 2 X X   | · · · · <b>r</b> · · · ·                     |

| ORF | Product name |
|-----|--------------|
|-----|--------------|

#### Alginate

PA2867probable chemotaxis transducerPA3819conserved hypothetical proteinPA5182hypothetical protein

#### RsmA

PA0105 cytochrome c oxidase, subunit II PA0108 cytochrome c oxidase, subunit III PA0129 gamma-aminobutyrate permease PA0173 probable methylesterase PA0210 malonate decarboxylase delta subunit malonate decarboxylase gamma subunit PA0212 PA0235 4-hydroxybenzoate transporter PcaK PA0396 twitching motility protein PilU PA0652 transcriptional regulator Vfr hypothetical protein PA0673 PA0706 chloramphenicol acetyltransferase PA0710 lactovlglutathione lvase coat protein B of bacteriophage Pf1) PA0723 PA0843 phospholipase accessory protein PlcR precursor PA0844 hemolytic phospholipase C precursor PA0905 RsmA, regulator of secondary metabolites PA0994 usher CupC3 PA1001 anthranilate synthase component I PA1092 flagellin type B PA1148 exotoxin A precursor PA1151 pyocin S2 immunity protein PA1173 cytochrome c-type protein NapB precursor PA1247 alkaline protease secretion protein AprE PA1251 probable chemotaxis transducer gamma-glutamyltranspeptidase precursor PA1338 adenosine 5'-phosphosulfate (APS) kinase PA1393 PA1409 acetylpolyamine aminohydrolase PA1635 potassium-transporting ATPase, C chain translocation protein in type III secretion PA1695 PA1758 para-aminobenzoate synthase component I PA1770 phosphoenolpyruvate synthase PA1781 assimilatory nitrite reductase large subunit PA1947 ribose transport protein RbsA ribose operon repressor RbsR PA1949 PA1983 cytochrome c550 PA2007 maleylacetoacetate isomerase PA2009 homogentisate 1,2-dioxygenase PA2064 copper resistance protein B precursor chaperone CupA2 PA2129 PA2337 transcriptional regulator MtlR PA2493 RND multidrug efflux membrane fusion protein MexE precursor PA2494 RND multidrug efflux transporter MexF PA2495 Multidrug efflux outer membrane protein OprN precursor PA2509 muconate cycloisomerase I PA2926 histidine transport protein HisP PA3540 GDP-mannose 6-dehydrogenase AlgD PA3543 alginate biosynthetic protein AlgK precursor PA3544 Alginate production outer membrane protein AlgE precursor PA3549 alginate o-acetyltransferase AlgJ diacylglycerol kinase PA3603 PA4110 beta-lactamase precursor 2-oxo-hept-3-ene-1,7-dioate hydratase PA4127 PA4138 tyrosyl-tRNA synthetase ferric enterobactin transport protein FepC PA4158 PA4205 hypothetical protein PA4210 probable phenazine biosynthesis protein PA4307 chemotactic transducer PctC PA4310 chemotactic transducer PctB PA4356 xenobiotic reductase

Function class

Adaptation, Protection

Membrane proteins

Membrane proteins

Energy metabolism Energy metabolism Transport of small molecules Adaptation, Protection Carbon compound catabolism Carbon compound catabolism Membrane proteins Motility & Attachment Transcriptional regulators Hypothetical, unclassified, unknown Antibiotic resistance and susceptibility Central intermediary metabolism Related to phage, transposon, or plasmid Secreted Factors (toxins, enzymes, alginate) Secreted Factors (toxins, enzymes, alginate) Translation, post-translational modification, degradation Motility & Attachment Adaptation, Protection Motility & Attachment Secreted Factors (toxins, enzymes, alginate) Adaptation, Protection Energy metabolism Secreted Factors (toxins, enzymes, alginate) Adaptation, Protection Amino acid biosynthesis and metabolism Amino acid biosynthesis and metabolism Carbon compound catabolism Transport of small molecules Protein secretion/export apparatus Biosynthesis of cofactors, prosthetic groups and carriers Energy metabolism Central intermediary metabolism Transport of small molecules Carbon compound catabolism Energy metabolism Carbon compound catabolism Carbon compound catabolism Adaptation, Protection Motility & Attachment Transcriptional regulators Transport of small molecules Transport of small molecules Membrane proteins Carbon compound catabolism Amino acid biosynthesis and metabolism Cell wall / LPS / capsule Cell wall / LPS / capsule Membrane proteins Cell wall / LPS / capsule Fatty acid and phospholipid metabolism Adaptation, Protection Carbon compound catabolism Amino acid biosynthesis and metabolism Transport of small molecules Membrane proteins Secreted Factors (toxins, enzymes, alginate) Adaptation, Protection Adaptation, Protection Adaptation, Protection

| ORF    | Product name   | Function class  |
|--------|--|---|
|        |  |   |
| PA4522 | beta-lactamase expression regulator AmpD                 | Cell wall / LPS / capsule                                 |
| PA4527 | still frameshift type 4 fimbrial biogenesis protein PilC | Motility & Attachment                                     |
| PA4855 | phosphoribosylamineglycine ligase                        | Nucleotide biosynthesis and metabolism                    |
| PA4953 | chemotaxis protein MotB                                  | Membrane proteins   |
| PA4954 | chemotaxis protein MotA                                  | Adaptation, Protection                                    |
| PA5041 | type 4 fimbrial biogenesis protein PilP                  | Motility & Attachment                                     |
| PA5042 | type 4 fimbrial biogenesis protein PilO                  | Motility & Attachment                                     |
| PA5043 | type 4 fimbrial biogenesis protein PilN                  | Motility & Attachment                                     |
| PA5044 | type 4 fimbrial biogenesis protein PilM                  | Motility & Attachment                                     |
| PA5243 | delta-aminolevulinic acid dehydratase                    | Biosynthesis of cofactors, prosthetic groups and carriers |
| PA5355 | glycolate oxidase subunit GlcD                           | Central intermediary metabolism                           |
| PA5361 | two-component sensor PhoR                                | Two-component regulatory systems                          |
| PA5501 | permease of ABC zinc transporter ZnuB                    | Membrane proteins   |
|        |  | 1   |